

Efficient Synthesis and in Silico Studies of the Benzimidazole Hybrid Scaffold with the Quinolinyloxadiazole Skeleton with Potential α -Glucosidase Inhibitory, Anticoagulant, and Antiplatelet Activities for Type-II Diabetes Mellitus Management and Treating Thrombotic Disorders

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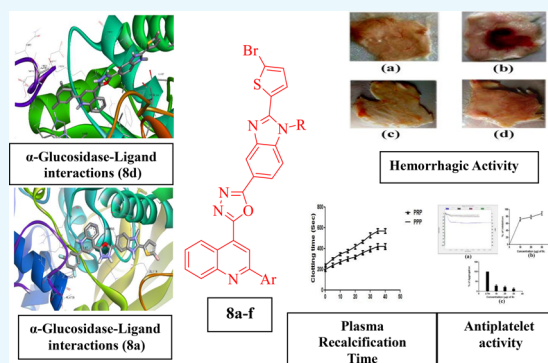
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Supporting Information

ABSTRACT: The current study evaluates antidiabetic, anticoagulant, and antiplatelet activity of novel benzimidazole-containing quinolinyloxadiazoles. These derivatives are synthesized and characterized using spectroscopy (FT-IR, ¹H NMR, and mass spectroscopy) and single-crystal X-ray diffraction methods. The inhibitory effects of these compounds were evaluated by the α -glucosidase inhibitory assay and shows the activity in the range of $IC_{50} = 0.66 \pm 0.05$ to 3.79 ± 0.46 μ g/mL. In addition, molecular docking studies revealed that benzimidazole-containing quinolinyloxadiazoles can correctly dock into the target receptor protein of the human intestinal α -glucosidase, while their bioavailability/drug-likeness was predicted to be acceptable but requires further optimization. On the other hand, compound **8a** and **8d** showed anticoagulant activity as they enhanced the clotting time from control 180–410 and 180–390 s, respectively, in platelet rich plasma and 230–460 and 230–545 s in platelet poor plasma. Furthermore, only **8a** showed antiplatelet activity by inhibiting epinephrine-induced platelet aggregation, and the observed aggregation inhibition was found to be 93.4%. Compounds **8a–f** show nontoxic properties because of the non-hydrolyzing properties in the RBC cells. In addition, **8a** and **8d** show anti-edema and anti-hemorrhagic properties in the experimental mice. These findings reveal that benzimidazole-containing quinolinyloxadiazoles act as α -glucosidase inhibitors to develop novel therapeutics for treating type-II diabetes mellitus and can act as lead molecules in drug discovery as potential antidiabetic and antithrombotic agents.



INTRODUCTION

As one of the chronic diseases, diabetes mellitus is expanding worldwide rapidly. It is characterized by high blood sugar levels for a longer period.¹ This uncontrolled hyperglycemia can cause serious damage to many vital organs in the body, including the kidneys, heart, and nervous tissues.^{2,3} Postprandial hyperglycemia has emerged as a prominent and early defect in type-2 diabetes and as a predictor of cardiovascular or even all-cause mortality, as well as an independent risk factor for atherosclerosis.⁴

The membrane-bound enzyme, α glucosidase, is found in the epithelium of the small intestine. It cleaves α -1,4 glycosidic

linkage at the nonreducing end of starch and disaccharides to release glucose units. Diabetes is caused with increasing glucose levels in the blood stream and relatively increases postprandial blood glucose levels.^{5–7} Inhibition of α -glucosidase is involved in the reduction of glucose absorption (rate) in the intestine and further decreasing plasma glucose levels. With these properties, α -glucosidase (therapeutic target) is used for the modulation of postprandial hyperglycemia in

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type-2 diabetes.⁸ The inhibitors, acarbose, miglitol, and voglibose, used to inhibit α -glucosidase are used to control postprandial blood glucose levels in type-2 diabetic patients.^{9,10}

Over the past decade, α -glucosidase inhibitors, such as acarbose, miglitol, and voglibose, have been used to decrease the postprandial blood glucose levels in type-2 diabetic patients.^{9,10} Furthermore, various studies show that the molecules which exhibit the α -glucosidase inhibitor property are also used in treating cancer, HIV, virus, and tumors.^{11–13} Therefore, the present scenario needs a proper designing and synthesizing of better α -glucosidase inhibitors to enrich the discipline of medicinal chemistry.

Blood plays a pivotal role in supplying micronutrients and macronutrients to different parts of the body.¹⁴ Meanwhile, blood oozing out during a vascular injury leads to many consequences. Thus, prevention of blood loss plays a major role in physiological condition and the phenomenon is termed as hemostasis, which is a highly regulated pathway.¹⁵ Some environmental and genetic factors may alter the hemostatic pathway, which leads to thrombosis.¹⁶ Thrombosis is nothing but the generation of unusual clots in arteries and veins, which is a major cause for death around the world.¹⁷ Whereas, antiplatelet and anticoagulant agents play a major role in treating thrombotic disorders.¹⁸ The present study finds the beneficial role of benzimidazole-containing quinolinyl oxadiazoles on thrombotic disorders.

Several quinoline derivatives show various pharmacological activities such as antifungal,^{19,20} antimalarial,²¹ antibacterial,²² anthelmintic,^{23,24} anticancer,^{25,26} anticonvulsant,²⁷ anti-inflammatory,²⁸ analgesic,²⁹ and antihyperglycemic activities.^{30–32} The quinoline derivatives isolated from natural systems have been reported for potent α -glucosidase inhibition.^{33–36}

On the other hand, benzimidazoles have potency for the anti-inflammatory, antispasmodic, analgesic, antihistaminic, antidiabetic, antimicrobial, antitubercular, antiproliferative, antitumor, anti-HIV-RT, anticancer, antiulcer, and cyclooxygenase inhibitor activities.^{37–46} The benzimidazole derivatives also exhibited α -glucosidase^{47–49} as well as antidiabetic⁵⁰ activity; hence, it is important to obtain novel benzimidazole derivatives as antidiabetic compounds. There are only limited studies of benzimidazole derivatives used on α -glucosidase inhibitory (AGI) activity and cytotoxicity.⁵¹

It was reported that oxadiazoles act as a potential class of α -glucosidase inhibitors^{54,52,53} and are found to have pharmacological activities such as anticancer,^{54,55} antimicrobial,^{56–58} anti-inflammatory,⁵⁹ anticonvulsant,⁶⁰ antioxidant,⁶¹ and antidiabetic.^{34,62} The recent report on 1,3,4-oxadiazole scaffolds discusses potential antiproliferative agents.^{63–65}

Drugs in market containing quinoline, benzimidazole, and 1,3,4-oxadiazole as a heterocycle shown in the Figure 1.

In view of the above observations in the quest to design better drugs for diabetes and thrombotic disorders in our current work, we have synthesized benzimidazole-containing quinolinyl oxadiazoles and subjected to anticoagulant, antiplatelet, and α -glucosidase inhibition activities.

RESULTS AND DISCUSSION

Chemistry. The synthetic methods for benzimidazole-containing quinolinyl oxadiazoles **8a–f** are illustrated in Scheme 1. The synthesis of precursors substituted quinoline-4-carboxylic acids **3a–b** with various acetophenones, as shown in Scheme 2. Furthermore, key intermediates **7a–c** were obtained initially from the core nucleus benzimidazole-5-

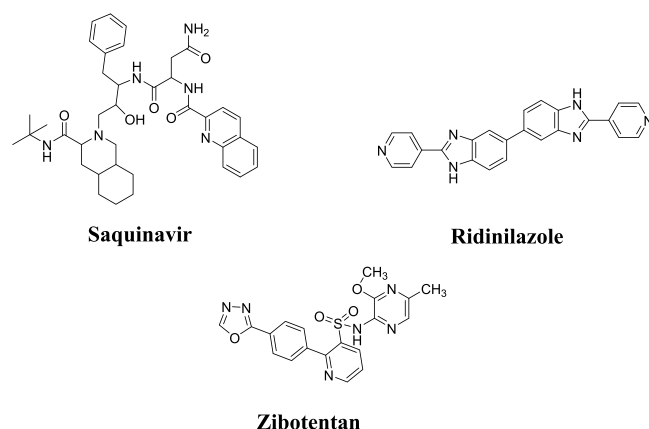


Figure 1. Drugs in market containing quinoline, benzimidazole, and 1,3,4-oxadiazole moieties.

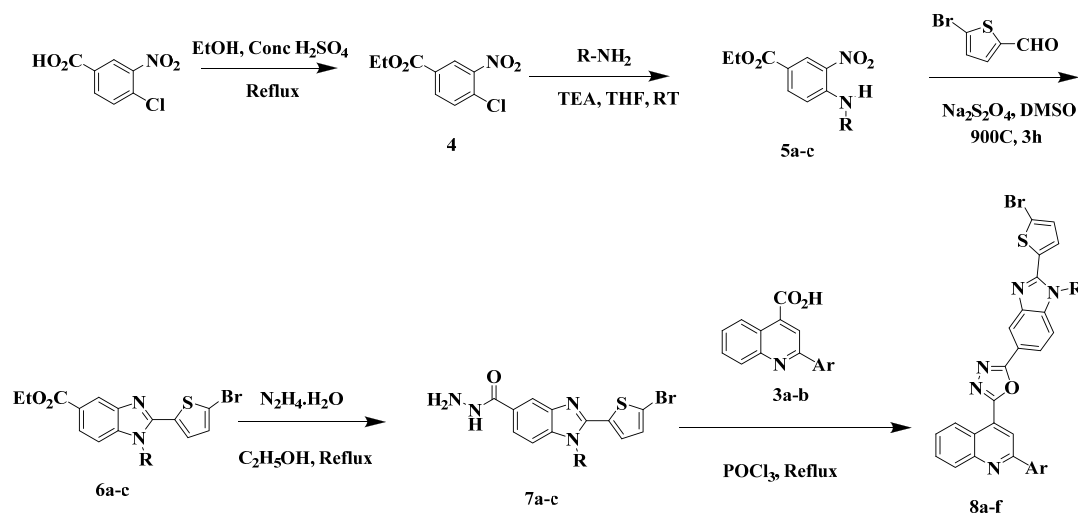
carboxylates **6a–c** that were efficiently synthesized by the “one pot” nitro reductive cyclization reaction between ethyl 3-nitro-4-(substituted amino) benzoates **5a–c** and 5-bromothiophene-2-carbaldehyde using sodium dithionite in dimethylsulfoxide. Interestingly, the “one-pot” reaction was proceeded very smoothly, in short reaction time with an excellent yield. The benzimidazole esters **6a–c** were converted into the corresponding hydrazides **7a–c** and then to highly functionalized quinolinylloxadiazoles **8a–f**. All the studied compounds were synthesized by using the same procedure and characterized by spectroscopic analysis.^{66,67} The solid-state properties of the target compounds are listed in Supporting Information Table S1.

X-ray Diffraction Analysis and Hirshfeld Surfaces Analysis. Figure 2 describes the ORTEP of the **6c**. The dihedral angle between rings Cg1: S1/C11–C14 and Cg4: N1/C4–C10/N2 is 27.4(2)°. Crystal structure is stabilized through C1···H1B···O2 intermolecular hydrogen bond in the form of 1D-infinite chain and C16···H16A···Cg3 (C4–C9) intermolecular interactions (Figure 3 and Table 1). The molecular arrangement within the crystal structure is shown along *a*, *b*, and *c*-axis (Figure 3). The crystal data and refinement parameters are given in Table 2.

The intercontacts in the crystal structure are analyzed and visualized using Hirshfeld surfaces and 2D fingerprint plots (Figures 4 and 5). The intercontacts found for **6c** are Br···C—1.5%, Br···H—11.1%, Br···N—2.0%, Br···S—1.7%, C···C—1.8%, C···H—18.4%, H···H—39.3%, N···C—1.0%, N···H—4.9%, O···C—1.7%, O···H—7.9%, O···O—0.3%, S···C—0.4%, and S···H—8.2%. The 2D finger print plots of these intercontacts are shown in Figure 5. The major contributions are from H···H, C···H, O···H, and N···H when compared to other intercontacts.

Computational Screening (Structure–Activity Relationship). The docking study was implemented using AutoDock 4.2.6 software for benzimidazole-containing quinolinyl oxadiazole derivatives and acarbose against N-terminal domain of the human intestinal α -glucosidase. The parameters of binding energy and binding mode of each compound were carefully analyzed. Using these parameters, hydrogen bond interactions and intermolecular interactions (π – π) with the enzymes were identified. Docking studies corroborate the significant interactions of all the in vitro active benzimidazole-containing quinolinylloxadiazole derivatives with the active site (acarbose inhibition site) of α -glucosidase enzyme. The

Scheme 1. Synthetic Route for the Preparation of 8a–f. Ar = 3,5-F₂C₆H₃, 2,4-Cl₂C₆H₃, R = CH₃, C₃H₇, C₄H₉



Scheme 2. Synthetic Route for the Preparation of 3a–b. Ar = 3,5-F₂C₆H₃, 2,4-Cl₂C₆H₃

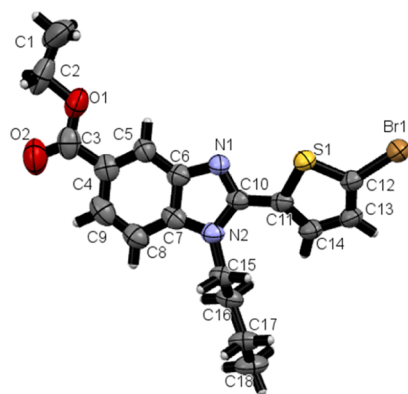
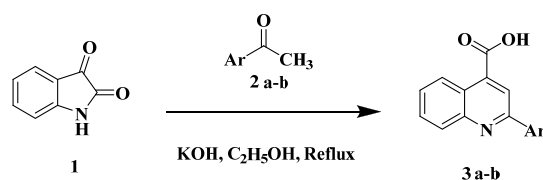


Figure 2. ORTEP diagram of the molecule 6c with thermal ellipsoids drawn at 50% probability.

docking scores also show positive correlation with the experimental results, in general. The docking pose of compounds 8a–f was shown in (Figure 6 and Table 3). The docking study predicted that the quinoline ring of compound 8d binds to enzyme forming π - π stacking with the amino group of Lys776 and benzimidazole ring forms π - π interaction with the amino group of Lys513 amino acid residue with the least binding energy of -8.45 kcal/mol. The second most active in this series is compound 8a with the least binding energy of -8.08 kcal/mol, in which quinoline ring forms π - π interaction with the amino group of Lys776 while benzimidazole ring forms π - π interaction with the amino

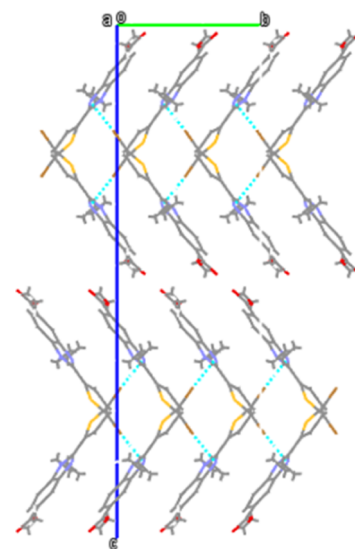


Figure 3. Molecular arrangement of the molecules viewed along the *b*-axis within the crystal structure. Dotted lines represent intermolecular hydrogen bonds.

group of Lys513. Furthermore, the benzimidazole ring of compound 8b forms π - π stacking with the amino group of Lys513 with the least binding energy of -7.77 kcal/mol, and in compound 8f with the least binding energy of -7.5 kcal/mol, the benzimidazole ring forms π - π interaction with amino group of Lys513. At the binding site of α -glucosidase, the amino groups of Lys776 and Lys513 residues construct the π - π stacking interaction with quinoline ring and benzimidazole ring of the ligand 8c, respectively. However, in the ligand 8e with binding energy -7.95 kcal/mol, no interaction has been seen with amino acid residues.

In general, all the ligand molecules except 8e showed more promising results against α -glucosidase compared to standard compounds. Hence, they may act as potential specific inhibitors for this target enzyme.

Biological Evaluation. Antidiabetic Activity: AGI Effect. α -Glucosidase inhibition assay was carried out for the newly synthesized compounds 8a–f in vitro. All the compounds

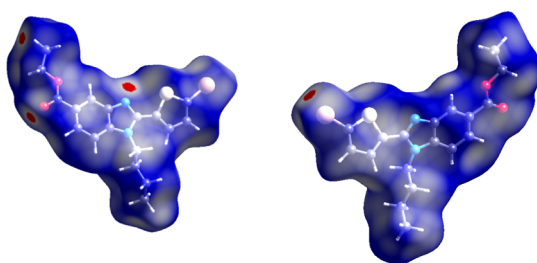
Table 1. Intramolecular and Intermolecular Hydrogen Bonds of Compound **6c**

D...H/X...A/Cg	D–H	H/X...A/Cg	D...A/Cg	D...H/X...A/Cg
C1...H1B...O2 ^a	0.96	2.54	3.412(13)	152
C16...H16A...Cg3 ^b		2.90	3.698(6)	141

$$^a -1/2 + x, 5/2 - y, 1 - z. \quad ^b 3/2 - x, -1/2 + y, z.$$

Table 2. Crystal Data and Refinement Statistics of Compound **6c**

formula	C ₁₈ H ₁₉ BrN ₂ O ₂ S
formula weight	407.32
crystal system	orthorhombic
space group	<i>Pbca</i> (no. 61)
<i>a</i> , <i>b</i> , <i>c</i> (Å)	11.1173(7), 9.6158(7), 34.457(2)
<i>V</i> (Å ³)	3683.5(4)
<i>Z</i>	8
<i>D</i> (calc) (g/cm ³)	1.469
μ (Mo <i>K</i> α) (/mm)	2.358
<i>F</i> (000)	1664
crystal size (mm)	0.21 × 0.23 × 0.26
Data Collection	
temperature (K)	293
radiation (Å)	Mo <i>K</i> α 0.71073
θ min–max (deg)	2.2, 26.4
dataset	–13: 13; –10: 12; –43: 43
tot., uniq. data, <i>R</i> (int)	370 38, 3761, 0.187
observed data [<i>I</i> > 0.0 σ (<i>I</i>)]	2377
Refinement	
<i>N</i> _{ref} , <i>N</i> _{par}	3761, 219
<i>R</i> , <i>wR</i> ₂ , <i>S</i>	0.0747, 0.2225, 1.05
max. and av. shift/error	0.03, 0.00
min. and max. resd. dens. (e/Å ³)	–0.63, 0.83

**Figure 4.** *d*_{norm} mapped on Hirshfeld surface for visualizing the intercontact in different orientations. Color scale is between –0.18 au (blue) to 1.4 au (red). The ball and stick model represents the molecule orientation.

exhibited significant α -glucosidase-inhibiting potential compared to standard acarbose ($IC_{50} = 1460.28 \pm 244.365 \mu\text{g}/\text{mL}$). It was demonstrated that compounds **8a** and **8d** had the best AGI activity with IC_{50} values 0.66 ± 0.05 and $0.68 \pm 0.02 \mu\text{g}/\text{mL}$ as represented in Table 4.

Plasma Recalcification Time. To pinpoint the possible role of **8a** and **8d** in the blood coagulation cascade, plasma recalcification time was accomplished using both human platelet rich plasma (PRP) and platelet poor plasma (PPP). Astonishingly, **8a** and **8d** displayed anticoagulant effect by enhancing the clotting time of both PRP and PPP from control 180–410, 180–390 and 230–460, 230–545 s, respectively. The supreme concentration consumed in both the cases was found to be 35 μg and remain unchanged upon increased dose to 40 μg , Figures 7 and 8. Coagulation factors are responsible

for blood clot which is a physiological phenomenon, and the blood coagulation cascade could be activated just to arrest the bleeding in the case of injury. It involves several factors such as factor VIIa to factor XIIIa, which mainly encompasses three pathways which are intrinsic, extrinsic, and common pathways. Some genetic imbalance and some environmental factors alter the normal coagulation system, which further leads to thrombosis, a pathological phenomenon. Plentiful of anticoagulants were reported from the biological sources and nanoparticles,^{14,68–71} but reports available on the anticoagulants from benzimidazole-containing quinolinyl oxadiazoles are very rare.

Antiplatelet and Nontoxic Properties of 8a–f. Furthermore, to ascertain the role of **8a** and **8d** on platelets, these were analyzed for platelet aggregation assay using PRP with agonist epinephrine. Curiously, only **8a** is able to inhibit the epinephrine-induced platelet aggregation but not the **8d** sample. The **8a** sample inhibits epinephrine-induced platelet aggregation of about 93% at the concentration of 30 μg as shown in Figure 9. However, platelets are very much essential to stop bleeding along with coagulation factors. Aggregation of platelets leads to the formation of platelet plugs when blood vessels are injured. Many physiological agents act as agonists such as ADP, epinephrine, collagen, thrombin, arachidonic acid, and thromboxane to activate platelets in order to arrest the blood by forming platelet plugs at the site injury. No control over the activation of platelets leads to pathophysiology of thrombotic disorders. Several antiplatelet agents have been identified from the biological sources,^{72,73} but very few are studied from synthesized compound sources.

Moreover, all the synthesized compounds of benzimidazole-containing quinolinyl oxadiazoles **8a–f** went unhydrolyzed with RBC cells. Molecules **8a** and **8d** were active in preventing hemorrhage and edema in experimental mice up to the concentration level of 100 μg . The positive control, *Daboia russelli* venom, encouraged hemorrhage and edema in experimental mice, suggesting its nontoxic property as shown in Figure 10.

CONCLUSIONS

Benzimidazole-containing quinolinyl oxadiazoles **8a–f** were synthesized. The novel synthesized compounds were characterized by spectral and analytical data and were screened for antidiabetic and antithrombotic activity. The results are correlated with docking studies. The molecular docking data provided positive correlation with in vitro antidiabetic activity in comparison with the standards revealed that these compounds can act as potential inhibitors. The core nucleus benzimidazole-5-carboxylates **6a–c** were efficiently synthesized by a “one pot” nitro reductive cyclization reaction between ethyl 3-nitro-4-(substituted amino) benzoates **5a–c** and 5-bromothiophene-2-carbaldehyde using sodium dithionite in dimethylsulfoxide. This “one-pot” reaction was proceeded very smoothly, in short reaction time with an excellent yield. Compounds benzimidazole-containing quinolinyl oxadiazoles **8a–f** showed good binding interactions with the target enzyme

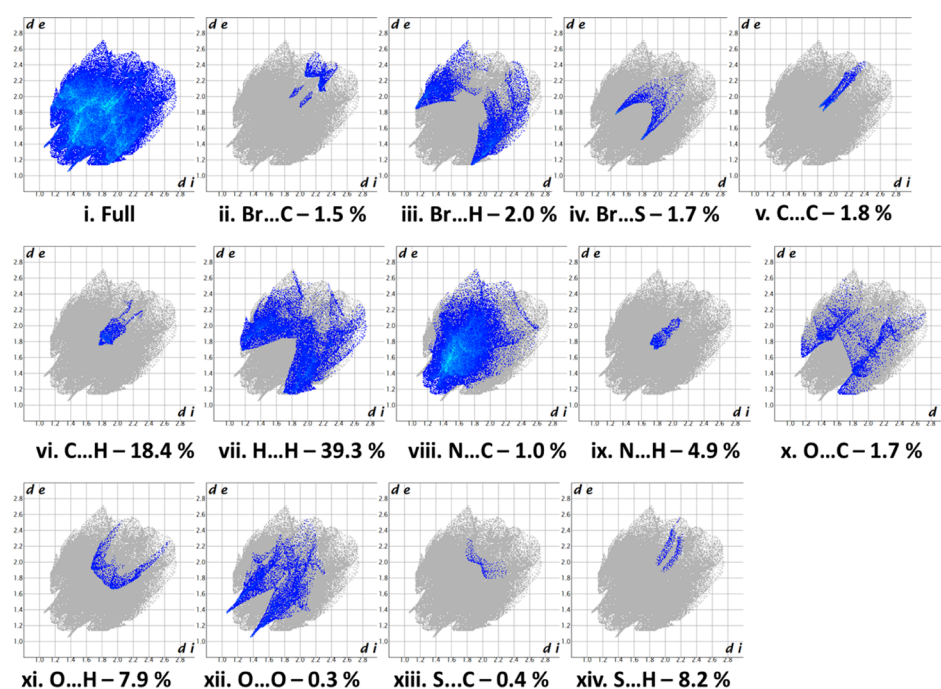


Figure 5. Fingerprint of the title compound 6c.

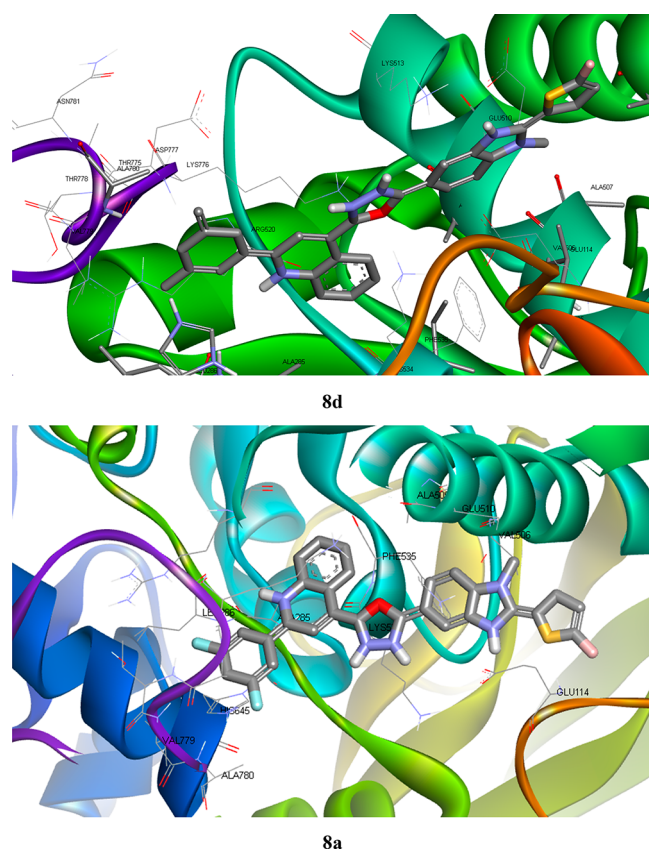


Figure 6. Docked poses of α -glucosidase–ligand interactions of 8d and 8a showing best binding affinity.

with least binding energies. Therefore, these compounds may be further evaluated and useful for diabetic treatment; on the other hand, this study for the first time demonstrates that benzimidazole-containing quinolinyl oxadiazoles (8a–f) are responsible for anticoagulant (8a and 8d) and antiplatelet (8a)

properties. Furthermore, they exhibit nontoxic property on RBC cells. Thus, they could play a major role in the treatment of thrombotic disorders, thus providing a new scope in drug discovery.

EXPERIMENTAL SECTION

Materials and Methods. All reagents were purchased from Sigma-Aldrich/Spectrochem, India, and used without further purification. Melting points were determined in an open capillary tube and were uncorrected. The progress of each reaction was monitored by ascending thin layer chromatography on silica gel G (Merck 1.05570.0001), visualized by UV light. The IR spectra for samples were recorded using a Shimadzu IRPrestige-21 FT-IR spectrophotometer, and the wave numbers were given in cm^{-1} . The ^1H NMR spectra were recorded ($\text{CDCl}_3/\text{DMSO}-d_6$ mixture) on an Agilent 400 MHz with an ATB probe and operated by VnmrJ software with TMS as the internal standard and Bruker; model AV 400. The X-ray intensity data were collected on a Rigaku Saturn70 diffractometer (Mo $K\alpha$ radiation) at 293 K, and data were collected and processed using Crystal Clear. Mass spectra were recorded in a WATERS model SynaptG2 LC-mass spectrometer.

Source for Anticoagulant and Antiplatelet Activity. Fresh human blood was collected from healthy donors for the PRP and PPP.

Preparation of 2-Arylquinoline-4-carboxylic Acids (3a–b). 2-arylquinoline-4-carboxylic acid (Figure 1) was prepared according to the literature method.^{74,75}

2-(3,5-Difluorophenyl)quinoline-4-carboxylic acid (3a): mp 230.0–232 °C.

2-(3,5-Dichlorophenyl)quinoline-4-carboxylic acid (3b): mp 198–204 °C.

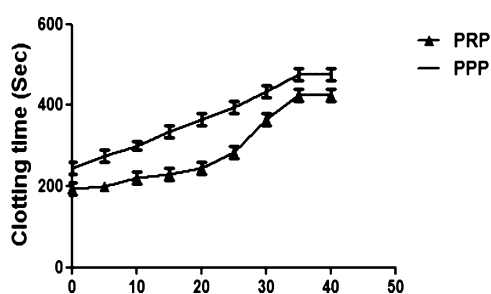
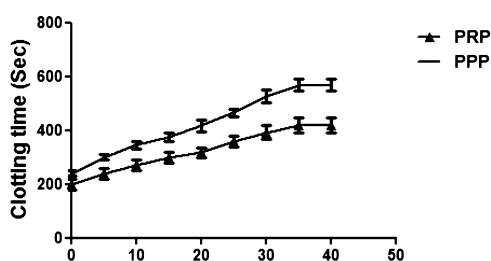
Preparation of Ethyl 4-(Alkylamino)-3-nitrobenzoates (5a–c). Ethyl 4-chloro-3-nitro benzoate (0.01 mol) was taken in 10 mL of tetrahydrofuran. To this solution, primary amine (0.015 mol) and triethylamine (0.03 mol) were

Table 3. Comparison of Predicted Binding Affinities of Active Benzimidazole-Containing Quinolinyl Oxadiazoles and Quinoline Schiff Bases and Acarbose against AGI Activity

ligand	binding energy	intermol. or internal energy	internal energy	torsional energy	unbound energy	H bonds/AA	π - π bonds/AA
8d	-8.45	-8.75	0	0.3	0	NA	2/Lys776, Lys513
8a	-8.08	-8.38	0.05	0.3	0.05	NA	2/Lys776, Lys513
8e	-7.95	-8.84	-0.46	0.89	-0.46	NA	NA
8b	-7.77	-8.66	-0.65	0.89	-0.65	NA	1/Lys513
8f	-7.5	-8.4	-0.51	0.89	-0.51	NA	1/Lys513
8c	-7.08	-7.98	-0.53	0.89	-0.53	NA, 3/Arg520, Lys534	2/Lys776, Lys513
acarbose	-14.38	-18.25	-5.52	3.88	-5.52	Val779	NA

Table 4. IC₅₀ Values of Benzimidazole-Containing Quinolinyl Oxadiazoles and Quinoline Schiff Bases for AGI Activity

sample	IC ₅₀ value ($\mu\text{g/mL}$) ($n = 3 \pm$ standard deviation)
8a	0.66 \pm 0.05
8b	2.76 \pm 0.37
8c	3.79 \pm 0.46
8d	0.68 \pm 0.02
8e	1.40 \pm 0.23
8f	2.81 \pm 0.24
acarbose	1460.28 \pm 244.365

**Figure 7.** Plasma recalcification time. 8a (0–40 μg) was pre-incubated with 0.2 mL of citrated human plasma PRP/PPP in the presence of 20 μL of 10 mM Tris-HCl buffer (pH 7.4) for 1 min at 37 $^{\circ}\text{C}$. CaCl_2 (20 μL ; 0.25 M) was added to the pre-incubated mixture and clotting time was recorded.**Figure 8.** Plasma recalcification time. 8d (0–40 μg) was pre-incubated with 0.2 mL of citrated human plasma PRP/PPP in the presence of 20 μL of 10 mM Tris-HCl buffer (pH 7.4) for 1 min at 37 $^{\circ}\text{C}$. CaCl_2 (20 μL ; 0.25 M) was added to the pre-incubated mixture and clotting time was recorded.

added. The reaction mixture was kept for stirring overnight at room temperature. After completion of the reaction, the reaction mixture was poured onto crushed ice. The product obtained was filtered, dried, and purified by recrystallization. Different ethyl 4-(alkylamino)-3-nitrobenzoates prepared using this method are as follows.

Ethyl 4-(Methylamino)-3-nitrobenzoate (**5a**): mp 98–100 $^{\circ}\text{C}$ (Lit. mp 101 $^{\circ}\text{C}$).⁷⁶

Ethyl 4-(Propylamino)-3-nitrobenzoate (**5b**): mp 77–78 $^{\circ}\text{C}$ (Lit. mp 78 $^{\circ}\text{C}$).⁷⁷

Ethyl 4-(Butylamino)-3-nitrobenzoate (**5c**): mp 52–54 $^{\circ}\text{C}$ (Lit. mp 50–52 $^{\circ}\text{C}$).⁷⁸

General Procedure for the Preparation of Ethyl 2-(5-Bromothiophen-2-yl)-1-alkyl-1H-benzo[d]imidazole-5-carboxylates (6a–c). Sodium dithionite (3.0 equiv) was added to a stirred solution of ethyl-4-(4-alkylamino)-3-nitrobenzoate (0.01 mol) and 5-bromothiophene-2-carbaldehyde (0.01 mol) in DMSO (20 mL). The reaction mixture was stirred at 90 $^{\circ}\text{C}$ for 3 h. After the completion of reaction, it was poured onto crushed ice. The solid separated was filtered off, washed with water, and dried. The crude product was recrystallized using dimethylformamide.

Ethyl 2-(5-Bromothiophen-2-yl)-1-methyl-1H-benzo[d]imidazole-5-carboxylate (6a). Yield 98%; mp 160–165 $^{\circ}\text{C}$; FT IR (ATR, ν cm^{-1}): 3040 (Ar-H), 2943 (C-H), 1713 (C=O), 1616 (C=N), 1538 (C=C); ^1H NMR (DMSO- d_6 δ ppm): 1.43 (t, 3H, CH_3), 4.04 (s, 3H, CH_3), 4.52 (q, 2H, CH_2), 7.05 (d, 1H, Br-thienyl-H, $J = 7.6$ Hz), 7.28 (d, 1H, Br-thienyl-H, $J = 7.6$ Hz), 7.48 (d, 1H, benzimidazolyl-C₆-H, $J = 8.4$ Hz), 7.66 (d, 1H, benzimidazolyl-C₇-H, $J = 8.4$ Hz), 8.03 (s, 1H, benzimidazolyl-C₄-H); ^{13}C NMR (100 MHz, CDCl_3 δ ppm): 14.4 (CH_3), 30.1 (CH_3), 61.5 (CH_2), 110.2, 110.5, 113.0, 129.5, 129.8, 130.9, 134.2, 135.1, 139.4, 141.5, 153.9, 167.2 (C=O); mass m/z : 364.54 ($\text{M} + \text{H}^+$), 366.79 ($\text{M} + \text{H} + 2$)⁺; Anal. found (calcd) for $\text{C}_{15}\text{H}_{13}\text{BrN}_2\text{O}_2\text{S}$ (%): C, 49.17 (49.33); H, 3.45 (3.59); 7.86 (7.67).

Ethyl 2-(5-Bromothiophen-2-yl)-1-propyl-1H-benzo[d]imidazole-5-carboxylate (6b). Yield 98%; mp 120–124 $^{\circ}\text{C}$; FT IR (ATR, ν cm^{-1}): 3060 (Ar-H), 2970 (C-H), 1728 (C=O), 1629 (C=N), 1462 (C=C); ^1H NMR (DMSO- d_6 δ ppm): 1.08 (t, 3H, CH_3), 1.39 (t, 3H, CH_3), 2.08–2.19 (m, 2H, CH_2), 4.35 (q, 2H, CH_2), 4.42 (t, 2H, CH_2), 7.13 (d, 1H, Br-thienyl-H, $J = 7.6$ Hz), 7.35 (d, 1H, Br-thienyl-H, $J = 7.6$ Hz), 7.55 (d, 1H, benzimidazolyl-C₆-H, $J = 8.4$ Hz), 7.71 (d, 1H, benzimidazolyl-C₇-H, $J = 8.4$ Hz), 8.09 (s, 1H, benzimidazolyl-C₄-H); ^{13}C (100 MHz, CDCl_3 δ ppm): 11.5 (CH_3), 14.1 (CH_3), 20.9 (CH_2), 49.6 (CH_2), 61.2 (CH_2), 111.2, 11.5, 114.8, 128.3, 129.2, 131.2, 135.4, 137.9, 139.2, 143.2, 154.2, 168.1 (C=O); mass m/z : 393.29 ($\text{M} + \text{H}^+$), 395.72 ($\text{M} + \text{H} + 2$)⁺; Anal. found (calcd) for $\text{C}_{17}\text{H}_{17}\text{BrN}_2\text{O}_2\text{S}$ (%): C, 51.75 (51.92); H, 4.17 (4.36); 7.30 (7.12).

Ethyl 2-(5-Bromothiophen-2-yl)-1-butyl-1H-benzo[d]imidazole-5-carboxylate (6c). Yield 98%; mp 95–98 $^{\circ}\text{C}$; FT IR (ATR, ν cm^{-1}): 3090 (Ar-H), 2985 (C-H), 1735 (C=O), 1612 (C=N), 1563 (C=C); ^1H NMR (DMSO- d_6 δ ppm): 0.96 (t, 3H, CH_3), 1.31 (t, 3H, CH_3), 1.39–1.51 (m, 2H, CH_2), 1.81–1.95 (m, 2H, CH_2), 4.22 (q, 2H, CH_2), 4.39 (t, 2H, CH_2), 7.19 (d, 1H, Br-thienyl-H, $J = 7.6$ Hz), 7.45 (d, 1H, Br-thienyl-H, $J = 7.6$ Hz), 7.56 (d, 1H, benzimidazolyl-C₆-

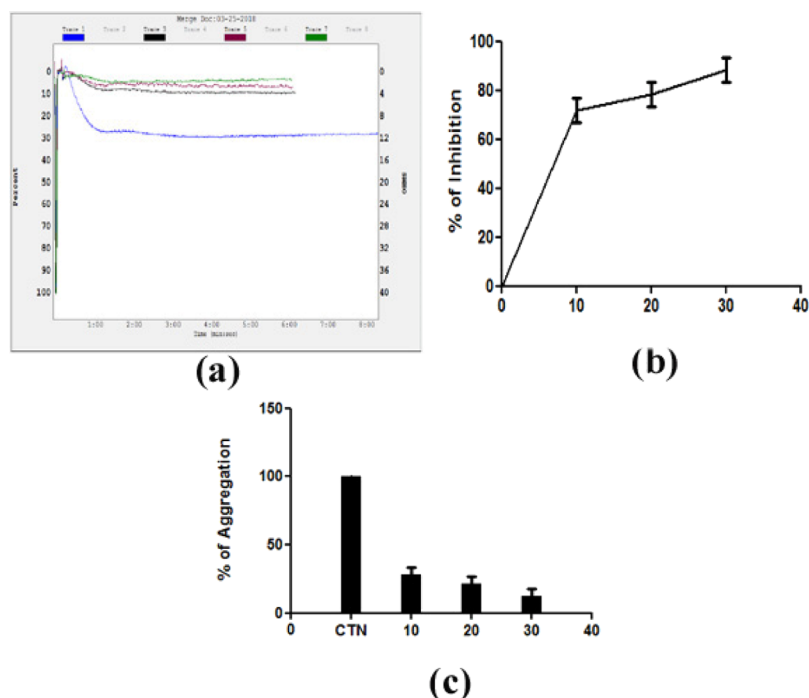


Figure 9. Platelet aggregation was initiated by adding epinephrine as an agonist of **8a**. (a) Traces of platelet aggregation: trace 1 (epinephrine 5 μM); trace 2 (epinephrine 5 μM + 10 μg of **8a**); trace 3 (epinephrine 5 μM + 20 μg of **8a**); and trace 4 (epinephrine 5 μM + 30 μg of **8a**). The values represent of three independent experiments. (b) Dose-dependent platelet aggregation inhibition %. (c) Dose-dependent platelet aggregation %.

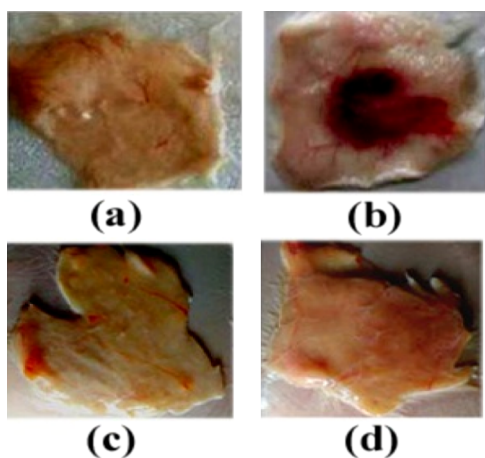


Figure 10. Dose-dependent hemorrhagic activity of **8a** and **8d**: (a) saline, (b) positive control 2 MDH venom, (c) 100 μg of **8a**, and (d) 100 μg of **8d** were injected independently into mice in a total volume of 50 μL intradermal.

H, $J = 8.4$ Hz), 7.79 (d, 1H, benzimidazolyl- $\text{C}_7\text{-H}$, $J = 8.4$ Hz), 8.12 (s, 1H, benzimidazolyl- $\text{C}_4\text{-H}$); ^{13}C NMR (100 MHz, CDCl_3 δ ppm): 13.9 (CH_3), 14.1 (CH_3), 21.9 (CH_2), 30.1 (CH_2), 49.8 (CH_2), 62.1 (CH_2), 110.1, 112.2, 113.9, 127.9, 132.5, 136.4, 137.3, 138.4, 140.8, 145.2, 155.3, 168.0 ($\text{C}=\text{O}$); mass m/z : 407.21 ($\text{M} + \text{H}$) $^+$, 409.53 ($\text{M} + \text{H} + 2$) $^+$; Anal. found (calcd) for $\text{C}_{18}\text{H}_{19}\text{BrN}_2\text{O}_2\text{S}$ (%): C, 53.25 (53.08); H, 4.55 (4.70); 6.71 (6.88).

General Procedure for the Preparation of 2-(5-Bromothiophen-2-yl)-1-alkyl-1H-benzo[d]imidazole-5-carbohydrazide (7a–c). A mixture of ethyl 2-(5-bromothiophen-2-yl)-1-alkyl-1H-benzo[d]imidazole-5-carboxylate (0.1 mol), hydrazine hydrate (0.15 mol), and 20 mL of ethanol

was refluxed on an oil bath for 10 h. The excess solvent was then distilled off under reduced pressure, and the concentrated solution was quenched with ice cold water. The solid separated was filtered, washed, and dried. The crude product was purified by recrystallization from ethanol.

2-(5-Bromothiophen-2-yl)-1-methyl-1H-benzo[d]imidazole-5-carbohydrazide (7a). Yield 98%; mp >230 $^\circ\text{C}$; FT IR (ATR, ν cm^{-1}): 3435, 3277 (NH_2), 3599 (NH), 3146 (Ar-H), 2947 (C-H), 1679 ($\text{C}=\text{O}$), 1663 ($\text{C}=\text{N}$), 1454 ($\text{C}=\text{C}$), 633 (C-S), 544 (C-Br); ^1H NMR ($\text{DMSO-}d_6$ δ ppm): 4.01 (s, 3H, CH_3), 4.79 (s, 2H, NH_2), 7.17 (d, 1H, Br-thienyl-H, $J = 7.6$ Hz), 7.40 (dd, 1H, benzimidazolyl- $\text{C}_6\text{-H}$, $J = 8.4$ Hz), 7.58 (dd, 1H, benzimidazolyl- $\text{C}_7\text{-H}$, $J = 8.4$ Hz), 7.67 (d, 1H, Br-thienyl-H, $J = 7.6$ Hz), 8.08 (s, 1H, benzimidazolyl- $\text{C}_4\text{-H}$), 9.96 (s, 1H, NH); ^{13}C NMR (100 MHz, CDCl_3 δ ppm): 30.8 (CH_3), 108.8, 113.0, 120.4, 128.7, 129.2, 129.8, 134.6, 134.8, 135.7, 141.0, 152.9, 165.4; mass m/z : 350.23 ($\text{M} + \text{H}$) $^+$, 352.57 ($\text{M} + \text{H} + 2$) $^+$; Anal. found (calcd) for $\text{C}_{13}\text{H}_{11}\text{BrN}_4\text{OS}$ (%): C, 44.58 (44.46); H, 3.30 (3.16); 15.78 (15.95).

2-(5-Bromothiophen-2-yl)-1-propyl-1H-benzo[d]imidazole-5-carbohydrazide (7b). Yield 98%; mp 140–144 $^\circ\text{C}$; FT IR (ATR, ν cm^{-1}): 3325, 3284 (NH_2), 3511 (NH), 3188 (Ar-H), 2968 (C-H), 1636 ($\text{C}=\text{O}$), 1533 ($\text{C}=\text{N}$), 1472 ($\text{C}=\text{C}$), 697 (C-S), 561 (C-Br); ^1H NMR ($\text{DMSO-}d_6$ δ ppm): 1.06 (t, 3H, CH_3), 2.06–2.18 (m, 2H, CH_2), 4.31 (t, 2H, CH_2), 4.81 (s, 2H, NH_2), 7.14 (d, 1H, Br-thienyl-H, $J = 7.6$ Hz), 7.44 (dd, 1H, benzimidazolyl- $\text{C}_6\text{-H}$, $J = 8.4$ Hz), 7.55 (d, 1H, Br-thienyl-H, $J = 7.6$ Hz), 7.77 (dd, 1H, benzimidazolyl- $\text{C}_7\text{-H}$, $J = 8.4$ Hz), 8.08 (s, 1H, benzimidazolyl- $\text{C}_4\text{-H}$), 10.02 (s, 1H, NH); ^{13}C NMR (100 MHz, CDCl_3 δ ppm): 12.5 (CH_3), 21.9 (CH_2), 49.6 (CH_2), 110.8, 113.0, 122.4, 125.7, 127.8, 132.8, 134.8, 136.5, 141.0, 152.9, 155.3, 165.4; mass m/z : 379.37 ($\text{M} + \text{H}$) $^+$, 381.71 ($\text{M} + \text{H} + 2$) $^+$;

Anal. found (calcd) for $C_{15}H_{15}BrN_4OS$ (%): C, 47.67 (47.50); H, 3.81 (3.99); 14.65 (14.77).

2-(5-Bromothiophen-2-yl)-1-butyl-1H-benzo[d]imidazole-5-carbohydrazide (7c). Yield 98%; mp >230 °C; FT IR (ATR, ν cm^{-1}): 3345, 3298 (NH₂), 3523 (NH), 3159 (Ar-H), 3009 (C-H), 1675 (C=O), 1621 (C=N), 1459 (C=C), 679 (C-S), 578 (C-Br); ¹H NMR (DMSO-*d*₆ δ ppm): 0.91 (t, 3H, CH₃), 1.29–1.42 (m, 2H, CH₂), 1.85–1.98 (m, 2H, CH₂), 4.37 (t, 2H, CH₂), 4.87 (s, 2H, NH₂), 7.25 (d, 1H, Br-thienyl-H, *J* = 7.6 Hz), 7.52 (dd, 1H, benzimidazolyl-C₆-H, *J* = 8.4 Hz), 7.68 (d, 1H, Br-thienyl-H, *J* = 7.6 Hz), 7.72 (dd, 1H, benzimidazolyl-C₇-H, *J* = 8.4 Hz), 8.16 (s, 1H, benzimidazolyl-C₄-H), 10.14 (s, 1H, NH); ¹³C NMR (100 MHz, CDCl₃ δ ppm): 13.9 (CH₃), 20.1 (CH₂), 30.1 (CH₂), 49.6 (CH₂), 109.3, 111.5, 115.8, 121.6, 129.1, 132.5, 133.3, 137.5, 139.2, 144.1, 148.2, 165.1; mass *m/z*: 393.29 (M + H)⁺, 395.68 (M + H + 2)⁺; Anal. found (calcd) for $C_{16}H_{17}BrN_4OS$ (%): C, 48.71 (48.86); H, 4.20 (4.36); 14.44 (14.25).

General Procedure for the Preparation of 2-(2-(5-Bromothiophen-2-yl)-1-alkyl-1H-benzo[d]imidazol-5-yl)-5-(2-(3,5-difluorophenyl)quinolin-4-yl)-1,3,4-oxadiazole (8a–f).⁶⁶ A mixture of 2-(5-bromothiophen-2-yl)-1-alkyl-1H-benzo[d]imidazole-5-carbohydrazide (1 mmol), substituted quinoline-4-carboxylic acid (1 mmol) and phosphorus oxychloride (2.5 mmol) was refluxed (100–110 °C for 6 h) and the residue was quenched with ice cold water. The solid separated was processed (filtered, washed, dried, and recrystallized).

2-(2-(5-Bromothiophen-2-yl)-1-methyl-1H-benzo[d]imidazol-5-yl)-5-(2-(3,5-difluorophenyl)quinolin-4-yl)-1,3,4-oxadiazole (8a). Yield 82%; mp 214–218 °C; FT IR (ATR, ν cm^{-1}): 3080 (Ar-H), 2969 (C-H), 1604 (C=N), 1478 (C=C), 1101 (C-F), 692 (C-S); ¹H NMR (DMSO-*d*₆ δ ppm): 3.97 (s, 3H, CH₃), 7.38 (overlapped d, 1H, difluorophenyl-C₄-H), 7.49 (d, 1H, Br-thienyl-H, *J* = 8.0 Hz), 7.64 (d, 2H, difluorophenyl, *J* = 7.2 Hz), 7.75 (d, 1H, benzimidazolyl-C₆-H, *J* = 8.4 Hz), 7.82 (d, 1H, Br-thienyl-H, *J* = 8.0 Hz), 7.91–8.01 (m, 2H, quinoliny-H), 8.17 (d, 1H, benzimidazolyl-C₇-H, *J* = 8.4 Hz), 8.25 (s, 1H, benzimidazolyl-C₄-H), 8.37 (d, 1H, quinoliny-H, *J* = 8.4 Hz), 8.56 (s, 1H, quinoliny-H), 8.77 (d, 1H, quinoliny-H, *J* = 8.0 Hz); ¹³C NMR (100 MHz, CDCl₃ δ ppm): 30.5 (CH₃), 109.0, 111.8, 114.0, 116.0, 116.2, 117.4, 120.2, 121.3, 126.5, 127.0, 127.7, 128.5, 129.1, 130.2, 130.8, 131.0, 132.3, 135.8, 136.7, 137.7, 142.0, 146.5, 153.9, 159.4, 161.7, 161.9, 164.7, 166.2; mass *m/z*: 600.41 (M + H)⁺, 602.29 (M + H + 2)⁺; Anal. found (calcd) for $C_{29}H_{16}BrF_2N_5OS$ (%): C, 58.19 (58.01); H, 2.81 (2.69); 11.51 (11.66).

2-(2-(5-Bromothiophen-2-yl)-1-propyl-1H-benzo[d]imidazol-5-yl)-5-(2-(3,5-difluorophenyl)quinolin-4-yl)-1,3,4-oxadiazole (8b). Yield 81%; mp 189–193 °C; FT IR (ATR, ν cm^{-1}): 3077 (Ar-H), 2962 (C-H), 1606 (C=N), 1477 (C=C), 1089 (C-F), 606 (C-S); ¹H NMR (DMSO-*d*₆ δ ppm): 1.10 (t, 3H, CH₃), 1.99–2.05 (m, 2H, -CH₂), 4.48 (t, 2H, -CH₂), 6.92 (overlapped d, 1H, difluorophenyl-C₄-H), 7.22 (d, 1H, Br-thienyl-H, *J* = 7.6 Hz), 7.55 (d, 1H, Br-thienyl-H, *J* = 7.6 Hz), 7.65–7.85 (m, 2H, quinoliny-H), 8.15 (d, 1H, benzimidazolyl-C₇-H, *J* = 8.4 Hz), 8.22 (d, 1H, benzimidazolyl-C₆-H, *J* = 8.4 Hz), 8.33 (d, 2H, difluorophenyl-H, *J* = 8.0 Hz), 8.44 (s, 1H, quinoliny-H), 8.65 (s, 1H, benzimidazolyl-C₄-H), 8.73 (d, 1H, quinoliny-H, *J* = 8.4 Hz), 9.17 (d, 1H, quinoliny-H, *J* = 8.4 Hz); ¹³C NMR (100 MHz, CDCl₃ δ ppm): 12.5 (CH₃), 21.9 (CH₂), 49.6 (CH₂), 108.0, 110.5, 113.0, 117.9,

123.0, 125.5, 125.9, 128.4, 129.0, 130.4, 130.8, 131.7, 133.4, 137.8, 139.1, 140.2, 141.3, 144.5, 145.9, 148.3, 151.1, 153.5, 156.5, 158.6, 159.3, 160.3, 162.8, 163.1; mass *m/z*: 628.10 (M + H)⁺, 630.25 (M + H + 2)⁺; Anal. found (calcd) for $C_{31}H_{20}BrF_2N_5OS$ (%): C, 59.10 (59.24); H, 3.07 (3.21); 11.33 (11.14). HPLC purity at 254 nm, 95.2%.

2-(2-(5-Bromothiophen-2-yl)-1-butyl-1H-benzo[d]imidazol-5-yl)-5-(2-(3,5-difluorophenyl)quinolin-4-yl)-1,3,4-oxadiazole (8c). Yield 80%; mp 192–194 °C; FT IR (ATR, ν cm^{-1}): 3051 (Ar-H), 2943 (C-H), 1623 (C=N), 1491 (C=C), 1123 (C-F), 623 (C-S); ¹H NMR (DMSO-*d*₆ δ ppm): 0.96 (t, 3H, CH₃), 1.27–1.35 (m, 2H, -CH₂), 2.18–2.25 (m, 2H, -CH₂), 4.58 (t, 2H, -CH₂), 7.34 (overlapped d, 1H, difluorophenyl-C₄-H), 7.41 (d, 1H, Br-thienyl-H, *J* = 8.0 Hz), 7.69 (d, 2H, difluorophenyl, *J* = 7.2 Hz), 7.77 (d, 1H, benzimidazolyl-C₆-H, *J* = 8.4 Hz), 7.84 (d, 1H, Br-thienyl-H, *J* = 8.0 Hz), 7.94–8.09 (m, 2H, quinoliny-H), 8.20 (d, 1H, benzimidazolyl-C₇-H, *J* = 8.4 Hz), 8.29 (s, 1H, benzimidazolyl-C₄-H), 8.40 (d, 1H, quinoliny-H, *J* = 8.4 Hz), 8.51 (s, 1H, quinoliny-H), 8.80 (d, 1H, quinoliny-H, *J* = 8.0 Hz); ¹³C NMR (100 MHz, CDCl₃ δ ppm): 13.9 (CH₃), 20.1 (CH₂), 30.1 (CH₂), 49.6 (CH₂), 109.0, 111.1, 118.5, 124.6, 126.1, 126.5, 129.0, 130.1, 131.0, 131.4, 132.3, 134.0, 138.4, 140.7, 141.8, 142.9, 145.1, 146.5, 149.9, 152.7, 154.1, 157.1, 159.2, 160.9, 161.9, 163.4, 165.5, 165.9; Mass *m/z*: 642.21 (M + H)⁺, 644.39 (M + H + 2)⁺; Anal. found (calcd) for $C_{32}H_{22}BrF_2N_5O_2S$ (%): C, 59.97 (58.82); H, 3.33 (3.45); 10.78 (10.90).

2-(2-(5-Bromothiophen-2-yl)-1-methyl-1H-benzo[d]imidazol-5-yl)-5-(2-(2,4-dichlorophenyl)quinolin-4-yl)-1,3,4-oxadiazole (8d). Yield 84%; mp >225 °C; FT IR (ATR, ν cm^{-1}): 3092 (Ar-H), 2951 (C-H), 1655 (C=N), 1455 (C=C), 777 (C-Cl), 649 (C-S); ¹H NMR (DMSO-*d*₆ δ ppm): 4.08 (s, 3H, CH₃), 7.23 (d, 1H, Br-thienyl-H, *J* = 8.0 Hz), 7.29 (d, 1H, Br-thienyl-H, *J* = 8.0 Hz), 7.45 (overlapped d, 1H, dichlorophenyl-C₃-H), 7.55 (s, 1H, benzimidazolyl-C₄-H), 7.63 (dd, 2H, benzimidazolyl-C₆-H and C₇-H, *J* = 8.4 Hz), 7.77–7.92 (m, 4H, dichlorophenyl-H and quinoliny-H), 8.18 (s, 1H, quinoliny-H), 8.54 (d, 1H, quinoliny-H, *J* = 8.4 Hz), 8.75 (d, 1H, quinoliny-H, *J* = 8.4 Hz); ¹³C NMR (100 MHz, CDCl₃ δ ppm): 30.8 (CH₃), 105.0, 108.2, 111.6, 114.4, 115.5, 120.7, 121.2, 122.0, 122.7, 123.3, 123.7, 124.4, 124.6, 125.2, 126.0, 127.7, 128.4, 129.5, 130.5, 130.9, 136.2, 140.7, 147.1, 148.4, 149.3, 153.1, 159.3, 160.4; mass *m/z*: 631.53 (M + H)⁺, 633.18 (M + H + 2)⁺; Anal. found (calcd) for $C_{29}H_{16}BrCl_2N_5OS$ (%): C, 55.13 (55.00); H, 2.41 (2.55); 11.22 (11.06). HPLC purity at 254 nm, 95.5%.

2-(2-(5-Bromothiophen-2-yl)-1-propyl-1H-benzo[d]imidazol-5-yl)-5-(2-(2,4-dichlorophenyl)quinolin-4-yl)-1,3,4-oxadiazole (8e). Yield 83%; mp 172–176 °C; FT IR (ATR, ν cm^{-1}): 3061 (Ar-H), 2981 (C-H), 1673 (C=N), 1463 (C=C), 759 (C-Cl), 635 (C-S); ¹H NMR (DMSO-*d*₆ δ ppm): 1.08 (s, 3H, CH₃), 2.29–2.38 (m, 2H, -CH₂), 4.75–4.82 (t, 2H, -CH₂), 7.32 (d, 1H, Br-thienyl-H, *J* = 8.0 Hz), 7.40 (overlapped d, 1H, dichlorophenyl-C₃-H), 7.47–7.53 (m, 2H, dichlorophenyl-H), 7.65–7.79 (m, 2H, quinoliny-H), 7.91 (d, 1H, Br-thienyl-H, *J* = 8.0 Hz), 8.19 (dd, 2H, benzimidazolyl-C₆-H and C₇-H, *J* = 8.4 Hz), 8.28 (s, 1H, benzimidazolyl-C₄-H), 8.37 (s, 1H, quinoliny-H), 8.42 (d, 1H, quinoliny-H, *J* = 8.4 Hz), 8.78 (d, 1H, quinoliny-H, *J* = 8.4 Hz); ¹³C NMR (100 MHz, CDCl₃ δ ppm): 11.5 (CH₃), 20.9 (CH₂), 48.6 (CH₂), 109.8, 112.0, 115.4, 118.2, 119.3, 124.5, 125.0, 125.7, 126.5, 126.8, 127.1, 127.5, 128.2, 128.6, 128.8, 129.0, 129.7,

132.2, 133.3, 134.3, 134.7, 140.0, 142.5, 144.5, 151.9, 157.1, 162.2, 162.7; mass m/z : 659.75 ($M + H$)⁺, 661.29 ($M + H + 2$)⁺; Anal. found (calcd) for C₃₁H₂₀BrCl₂N₅OS (%): C, 56.17 (56.30); H, 3.21 (3.05); 10.44 (10.59).

2-(2-(5-Bromothiophen-2-yl)-1-butyl-1H-benzo[d]-imidazol-5-yl)-5-(2-(2,4-dichlorophenyl)quinolin-4-yl)-1,3,4-oxadiazole (8f). Yield 85%; mp >225 °C; FT IR (ATR, ν cm⁻¹): 3049 (Ar-H), 2933 (C-H), 1644 (C=N), 1444 (C=C), 741 (C-Cl), 658 (C-S); ¹H NMR (DMSO-*d*₆ δ ppm): 0.90 (s, 3H, CH₃), 1.33–1.40 (m, 2H, -CH₂), 2.81–2.89 (m, 2H, -CH₂), 4.95 (t, 2H, -CH₂), 7.28 (d, 1H, Br-thienyl-H, $J = 8.0$ Hz), 7.35 (overlapped d, 1H, dichlorophenyl-C₃-H), 7.49–7.58 (m, 2H, dichlorophenyl-H), 7.65–7.76 (m, 2H, quinolinyl-H), 7.88 (dd, 2H, benzimidazolyl-C₆-H and C₇-H, $J = 8.4$ Hz), 7.97 (d, 1H, Br-thienyl-H, $J = 8.0$ Hz), 8.24 (s, 1H, benzimidazolyl-C₄-H), 8.33 (s, 1H, quinolinyl-H), 8.45 (d, 1H, quinolinyl-H, $J = 8.4$ Hz), 8.81 (d, 1H, quinolinyl-H, $J = 8.4$ Hz); ¹³C NMR (100 MHz, CDCl₃ δ ppm): 14.5 (CH₃), 22.1 (CH₂), 33.1 (CH₂), 48.6 (CH₂), 112.2, 115.2, 118.6, 120.4, 121.5, 127.7, 128.2, 129.7, 130.0, 131.4, 131.6, 132.2, 133.0, 134.7, 135.5, 136.4, 137.5, 137.9, 144.2, 147.7, 154.1, 155.4, 156.3, 160.1, 165.2, 166.3, 167.4; mass m/z : 674.29 ($M + H$)⁺, 676.54 ($M + H + 2$)⁺; Anal. found (calcd) for C₃₂H₂₀BrCl₂N₅OS (%): C, 56.79 (56.91); H, 3.17 (3.28); 10.49 (10.37).

Single Crystal X-ray Diffraction Studies. The single-crystal X-ray diffraction intensities collection and data process were carried out using the Rigaku Saturn 724 diffractometer (Mo $K\alpha$ radiation; 296 K) and *CrystalClear* software, respectively.⁷⁹ Direct methods were used for solving the structure (SHELXS), and refinement was made by the full-matrix least squares method on F^2 using *SHELX*.⁸⁰ All the non-hydrogen atoms were exhibited in the first difference Fourier map itself, in which all the hydrogen bonds were geometrically positioned (C-H = 0.93 Å, O-H = 0.82 Å) and a riding model with $U_{iso}(H) = 1.2U_{eq}$ and $1.5U_{eq}(O)$ was used for the refinement. The geometry of the **6c** is calculated using *PLATON* program.⁸¹ The ORTEP and packing diagrams were generated using *MERCURY*.⁸² Table 2 represents the details of crystal structure and data refinement. Figure 2 represents the ORTEP⁸² of the molecule with thermal ellipsoids drawn at 50% probability.

Hirshfeld Surfaces Analysis. The Hirshfeld surfaces computational analysis was accomplished to determine the intercontacts in the crystal structure of compound **6c** as shown in Figure 4, in which the intercontacts of the intermolecular interaction were indicated by the red colored spots over the Hirshfeld surface. The short interatomic contacts emerged by the strong hydrogen bonds were indicated by the dark red spots on the d_{norm} surface, while the light red spots indicate the other intermolecular interactions. The intercontacts with respect to d_i and d_e were plotted using 2D-fingerprint plot as shown in the Figure 5.^{74,83–86}

Computational Screening by Molecular Docking Studies. Protein and Ligand Preparation. The N-terminal domain of α -glucosidase protein in the human intestine (3CTT) was obtained from PDB (<http://www.rcsb.org/pdb>). The bound ligands, heteroatoms, and water molecules were removed from each of the protein molecules. Polar hydrogens and Gasteiger charges were assigned after merging of nonpolar hydrogen atoms. After preprocessing of the crystal structure, it was saved in the PDBQT file format using ADT. Ligands were optimized and saved in SDF format using Maestro 2D Draw

(version 9.7). Using Marvin Sketch (ver 17.1.9.0; <http://www.chemaxon.com>), all the SDF formatted ligands were converted to the PDB format and subsequently to the PDBQT format in AutoDockTools (ADT)-1.5.6.

Molecular Docking Simulations. AutoDock 4.2.6 was used to carry out the docking simulations. A standard protocol for molecular docking was performed. To validate drug–target association, automated docking was performed with AutoDock 4.2.6.⁸⁷ The grid map was set on the whole protein by AutoGrid 4 with grid dimensions to their maximum values. Ten independent genetic algorithm runs were performed for each ligand. The resulted values differing by <2.0 Å in positional root-mean-square deviation show the most favorable free energy of binding. All torsions were allowed to rotate during docking. Docked poses were visualized considering a maximum cutoff distance of 3.0 Å for hydrogen bonds. Analysis and visualizations were carried out using MacPymol,⁸⁸ ADT, and PoseView.⁸⁹

Antidiabetic activity: AGI Effect. AGI assay was performed by the hydrolysis of chromogenic substrate *p*-nitrophenyl- α -D-glucopyranoside (PNPG) using a previously explained method elsewhere.⁹⁰ Newly synthesized compounds **8a–f** each were dissolved in DMSO to obtain 1 mg/mL stock solution. From 1 mg/mL stock solution, further dilution was made into 0.1, 0.005, and 0.001 mg/mL using 0.1 M phosphate buffer (pH 7) to find their IC₅₀ values. Briefly, 100 μ L of reaction mixture in a 96-well microplate containing 40 μ L (0.0015 units) of α glucosidase enzyme, 50 μ L of compounds, and 10 μ L of phosphate buffer (0.1 M) pH 7 was incubated at 37 °C for 20 min. After that, 50 μ L of substrate (20 mM, PNPG) was added. The reaction mixture was further incubated at 37 °C for 30 min, and immediately, 100 μ L of sodium carbonate (200 mM) solution was added for terminating the reaction. The reaction blank was prepared by replacing compounds and enzyme with buffer. Similarly, the control was prepared using buffer in the place of compound. Also, sample blanks were prepared for each sample replacing enzyme with buffer. The acarbose (50 mg) tablets were dissolved in buffer at 10 mg/mL concentration and used as a standard. The absorbance of *p*-nitrophenol released was quantified by reading at 405 nm using a ThermoScientific Multiscan plate reader. The percent inhibition of α glucosidase enzyme was calculated using the following formula.

$$\text{Inhibition (\%)} = \left[\frac{(\text{absorbance}_{\text{control}} - \text{absorbance}_{\text{sample}})}{\text{absorbance}_{\text{control}}} \right] \times 100$$

The amount of sample required for the 50% inhibition of glucosidase enzyme was represented by IC₅₀ values.

Plasma Recalcification Time. Plasma recalcification time was measured as described by Quick et al.⁹¹ Different concentrations of compounds **8a** and **8d** (0–40 μ g) were pre-incubated with 0.2 mL citrated human plasma at 37 °C for 1 min. Clotting was measured after the addition of 20 μ L CaCl₂ (0.25 M).

Preparation of PRP and PPP. PRP and PPP were prepared using the method described by Ardlie and Han.⁹² The concentration of platelets in PRP was conserved as 3.1×10^8 platelets/mL with PPP at 37 °C.

Platelet Aggregation.⁹³ Platelet aggregation was performed in a Chronology dual channel whole blood/optical lumi aggregation system (model-700) as described by Born et al.⁹³ PRP was pre-incubated with various concentrations of **6c** (0–

15 μg) in 0.25 mL reaction volume mixture. The platelet aggregation process was initiated by addition of agonists independently such as ADP and epinephrine.

Direct Hemolytic Activity. Direct hemolytic activity was performed as described previously.¹⁸ Concisely, packed human erythrocytes and phosphate-buffered saline (PBS) were mixed in the ratio 1:9. Different concentrations of compounds **8a–f** (100 μg) were pre-incubated with 1 mL of this above solution for 1 h at 37 °C. Then, the reaction was arrested by adding 9 mL of ice cold PBS and centrifuged at 1000g for 10 min at 37 °C. Finally, the amount of hemoglobin released was measured at 540 nm.

Edema Inducing Activity.⁹⁴ Edema inducing activity was performed as described by Sannanaik Vishwanath et al.⁹⁴ Briefly, compounds **8a** and **8b** were injected at different doses (10–100 μg) for the selected group of five mice at the right foot pads. After 1 h, mice were anaesthetized using diethyl ether, and then, the hind-limbs were removed carefully at the ankle joint and weighed. Saline (20 μL) was injected alone into the left foot serves as control. Minimum edema dose was defined as the amount of protein required to cause an edema ratio of 120%.

Hemorrhagic Activity.⁹⁵ Hemorrhagic activity was performed as described by Kondo et al.⁹⁵ Briefly, higher concentrations of compounds **8a** and **8d** were injected with saline into the selected group of mice. Mice that received only saline act as the negative control whereas a group of mice that received snake venom serves as the positive control. After 3 h, all groups of mice were anesthetized, and then, hemorrhage was noticed by removing the dorsal patch of skin surface carefully. The minimum hemorrhagic dose was defined as the amount of the compound producing 10 mm of hemorrhage in diameter.

Statistical Analysis. The data are presented as mean \pm SD. Statistical analyses were performed by Student's *t*-test. A significant difference between the groups was considered if $P < 0.01$.

■ ASSOCIATED CONTENT

● Supporting Information

The Supporting Information is available free of charge on the ACS Publications website at DOI: 10.1021/acsomega.8b01476.

Physical properties and structural features of target compounds; IR spectra, ¹H NMR spectra, RP-HPLC-spectrum, and LC-MS spectrum of specific compounds (PDF)

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Notes

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