






Molecular Modeling, Synthesis, and Antihyperglycemic Activity of the New Benzimidazole Derivatives – Imidazoline Receptor Agonists

Artur Martynov ¹, Boris Farber ², Tatyana Bomko ¹, Daniel L Beckles ³, Ilya Kleyn ⁴

¹Laboratory and Clinical department of Molecular Immunopharmacology, SI “ I. Mechnikov Institute of Microbiology and Immunology of National Academy of Medical Sciences of Ukraine, Kharkiv, Ukraine; ²R&D Department, Noigel LLC, New York, NY, USA; ³Baylor Scott & White Health, Round Rock, TX, USA; ⁴SUNY Downstate Medical Center / University Hospital of Brooklyn, New York, NY, USA

Correspondence: Artur Martynov, Email imiamn@gmail.com

Introduction: The paper presents the results of a study on the first synthesized benzimidazole derivatives obtained from labile nature carboxylic acids. The synthesis conditions of these substances were studied, their structure was proved, and some components were found to have sugar-reducing activity on the model of alloxan diabetes in rats.

Methods: The study used molecular modeling methods such as docking based on the evolutionary model (igemdock), RP_HPLC method to monitor the synthesis reaction, and 1H NMR and 13C NMR, and other methods of organic chemistry to confirm the structures of synthesized substances.

Results & Discussion: The docking showed that the ursodeoxycholic acid benzimidazole derivatives have high tropics to all imidazoline receptor carriers (PDB ID: 2XCG, 2bk3, 3p0c, 1QH4). The ursodeoxycholic acid benzimidazole derivative and arginine and histidine benzimidazole derivatives showed the highest sugar-lowering activity in the experiment on alloxan-diabetic rats. For these derivatives, the difference in glucose levels of treated rats was significant against untreated control. Therefore, the new derivatives of benzimidazole and labile natural organic acids can be used to create new classes of imidazoline receptor inhibitors for the treatment of diabetes mellitus and hypertension.

Keywords: imidazoline receptors, benzimidazole derivatives, synthesis, antihyperglycemic activity

Introduction

Research interest in imidazoline receptor (IR) agonists has recently increased.

This is because the selective I₁-imidazoline receptor (I₁R) agonist moxonidine,^{1,2} the selective I₂-imidazoline receptor (I₂R) agonists diakamph³ and metformin⁴ have been found to have the ability to stimulate telomerase expression in cells and exhibit the ability to inhibit senescence.

In addition, the apparent ability of diakamph (V), moxonidine (VI), and metformin to increase tissue sensitivity to insulin has been shown. All three substances have been successfully used as medicines. Metformin is used as a treatment for diabetes mellitus, and moxonidine (VI) is a centrally-acting hypotensive drug (Figure 1). Particular attention should be paid to I₃-imidazoline receptors (I₃R), which are found in the pancreas and stimulated, leading to increased insulin production by beta-cells.⁵

I₁Rs are associated with phosphatidylcholine hydrolysis, the formation of arachidonic acid and eicosanoids, and are actively associated with catecholamine synthesis. Some studies indicate that IIRs belong to neurocytokine receptors and are similar in action to interleukins.^{6–8} I₁Rs are located in the brainstem and modulate blood pressure. In peripheral tissues, I₁Rs are found in the kidneys and are associated with regulating sodium balance in the body.⁹ In the human brain, I₁Rs are concentrated in the triatum, pale globe, and substantia nigra.⁷

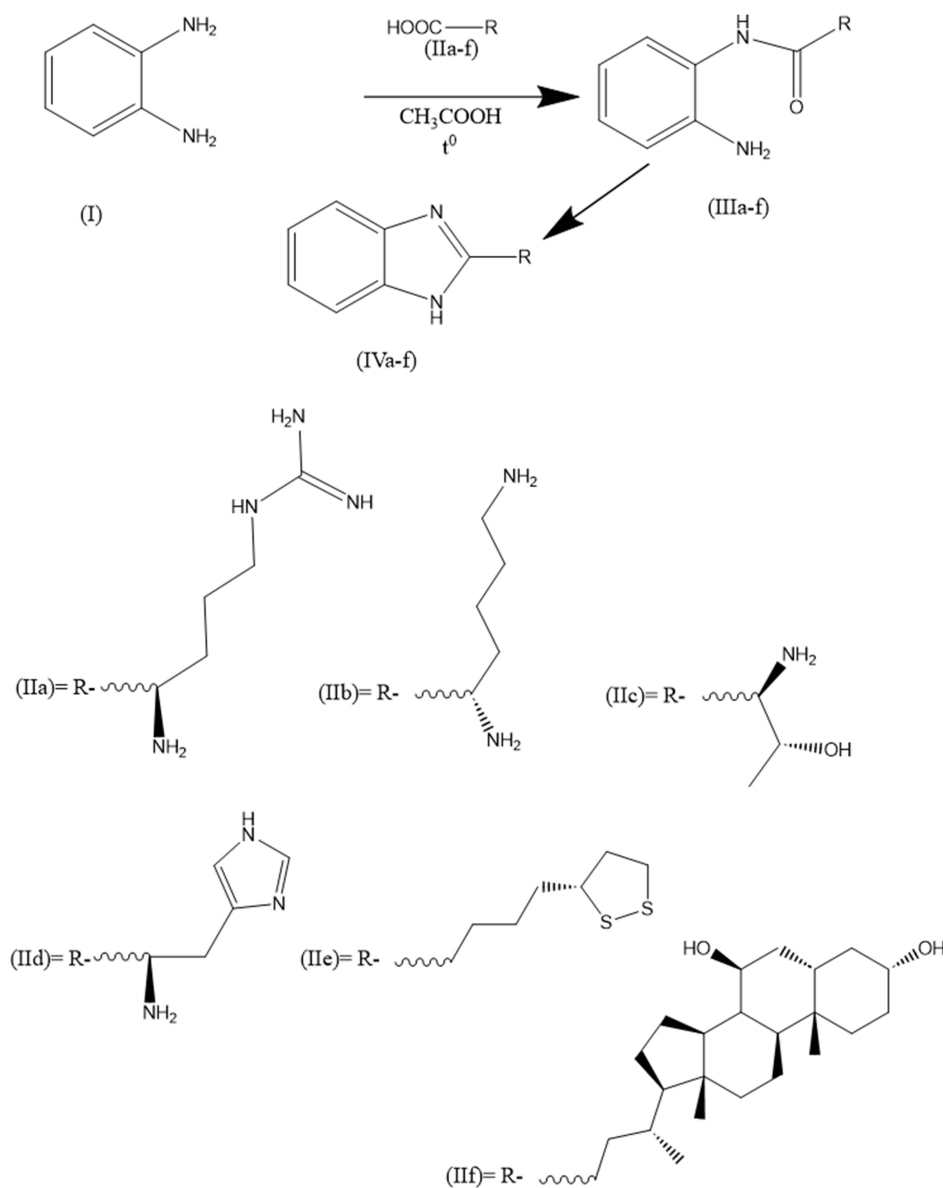


Figure 1 Synthesis of some benzimidazole derivatives (IVa-f).

The activation of I₁Rs also negatively regulates the progression of fibrosis in hepatic stellate cells through a Nrf2-dependent pathway. This hepatic I1-dependent antifibrotic action was also observed *in vivo* in mice.¹⁰

Imidazoline Subtype 1 receptors have been shown to affect insulin and adiponectin positively. A study using insulin-secreting Min6 b-cells found that activation of I1Rs with a selective I1 receptor ligand, S43126, had a glucose-dependent insulinotropic action.¹¹ This effect was prevented by efaroxan, a selective I₁Rs antagonist. Similarly, it was also demonstrated that activating I1Rs on insulin-targeted tissues such as liver and adipose tissues improved insulin sensitivity.¹² LNP599, a highly selective I₁Rs compound, stimulates adiponectin secretion in 3T3-L1 adipocyte cultures.¹³

In human platelets, dexmedetomidine, an alpha-2 adrenergic agonist and I1Rs agonist suppressed ADP-induced platelet aggregation when alpha-2 adrenergic receptors were blocked by yohimbine. This specific activation of I₁Rs led to an interesting antiaggregation effect on platelets.¹⁴

Selective I1 receptor agonists increased the expression of farnesoid X nuclear receptors in 3T3-L1 adipocyte cultures.¹⁵ This resulted in improved hepatic steatosis due to an increase in intracellular calcium and p38 phosphorylation. *In vivo*, an improvement in hepatic steatosis was observed in mice treated with rilmenidine.

I₂Rs are present both in the human brain and in peripheral tissues. The highest number of I₂Rs in the brain is found in the pineal gland.^{16,17} Decreased I₂R density in the human brain is associated with Huntington's chorea and Alzheimer's disease, as well as in drug addiction.⁷ I₂R was found most abundantly in mitochondrial membrane fractions derived from brain, kidney, heart, and liver.^{7,16} Some studies have found correlation of I₂R with monoamine oxidase (MAO).¹² In particular, purified I₂R protein has the same amino acid sequence with MAO-A and MAO-B, and imidazoline-containing substances inhibit MAO activity.¹⁷

Over the years, Idazoxan has been used to identify I₂Rs; however, it also binds to adrenoceptors. Because of this, medicinal chemistry has been actively developing selective I₂Rs ligands with low I₁ and α 2-adrenergic receptor binding activities. This has led to the development of a large library of selective I₂Rs ligands, and the effort continues today. For instance, a recent study describes a new family of (2-imidazolin-4-yl) phosphonates that have a high affinity for I₂Rs.¹⁸ Several compounds have been used in previous studies to explore the pharmacology of I₂Rs and have become valuable research tools. These include phenyzoline, 2-BFI, CR4056, BU224. Of note, two synthetic ligands have moved from preclinical research to human studies. One radiolabeled I₂Rs ligand, 11C-BU99008, has been used in PET studies with human volunteers to demonstrate receptor distribution in the brain.¹⁹ In the future, the PET ligand 11C-BU99 could help expand our understanding of I₂Rs in neurodegenerative disorders. This is because the density of I₂Rs changes in several psychiatric conditions, especially where there is marked gliosis. Therefore, the ligand could be used to confirm and monitor neurodegenerative diseases as well as detect glioblastomas. The authors of the study have previously established that 11C-BU99 has low affinity for MAOB but high affinity and selectivity for I₂Rs.²⁰

I₂Rs ligands have been studied extensively for their potential analgesic effects. These effects have been demonstrated by various compounds in multiple preclinical pain models. However, the research suggests that these compounds are not very effective in treating acute nociception in thermal stimulus-induced pain models. Previous studies have shown that I₂Rs ligands, including trazizoline, LSL601012, BFI, and phenyzoline, do not produce significant antinociception in rodent models of acute nociception, such as radiant tail flick, and hot plate assays.^{21–23} Li's group conducted recent studies that support these findings. Mild antinociception was observed with the use of 2-BFI and phenyzoline, while selective I₂ receptor ligands such as BU224 and S22687 (5-[2methyl phenoxy methyl]-1,3-oxazolin-2-yl) amine) did not have any effect on the warm water tail flick assay.^{24,25} I₂Rs agonists are generally effective in reducing pain caused by chemical stimuli. For instance, in a writhing test 2BFI, morphine and BU224 were found to have equal efficacy in reducing writhing response.²⁶ During the formalin test, rodents exhibit nocifensive behaviors that consist of two phases: an early neurogenic Phase I, followed by a later inflammatory Phase II.²⁷ During Phase II, CR4056, 2-BFI, and BU224 all demonstrated a dose-dependent reduction in the flinching response.²⁸ In a different study, a new I₂Rs agonist called CR4056 completely reversed capsaicin-induced neurogenic/inflammatory allodynia; the effect was dose-dependently blocked by idazoxan.²⁹

Several studies have shown that both 2-BFI and phenyzoline can reduce mechanical allodynia in rats with chronic constriction injury (CCI) peripheral neuropathic pain.²⁸ It has been confirmed that idazoxan attenuates the antiallodynic effects of 2-BFI and phenyzoline, which indicates that the mechanism is mediated by the I₂ receptor.^{28,30} In multiple rat models of chronic pain, CR4056 has shown a strong antihyperalgesic effect. It has been found to reduce mechanical allodynia in rats treated with complete Freund's adjuvant (CFA), attenuate mechanical hyperalgesia in diabetes-induced neuropathic pain, and reduce neuropathic pain induced by chronic treatment with the chemotherapeutic agent bortezomib.^{29,31} Additionally, CR4056 is effective in a rat model of fibromyalgia, reversing acidic saline-induced mechanical allodynia. In a postoperative pain model, CR4056 was fully effective in reversing mechanical allodynia, and this effect was blocked by idazoxan.³² Interestingly, no sex difference was found for CR4056-induced antinociception. These studies show that I₂Rs agonists can reduce chronic pain and may be a new form of painkillers with a wide range of effectiveness.

The I₃Rs are believed to be a separate binding site from I₁₋₂Rs. However, not much is known about their nature and function, despite previous research.^{33,34} Studies have shown that phentolamine and other imidazoline compounds can affect glycemic control in humans.³⁵ In pancreatic islet tissue, protein bands consistent with I₂Rs have been identified using an antiserum. However, the existence of I₁Rs is disputed. Research has shown that some I₂Rs agonists can regulate insulin secretion in pancreatic b-cells, while others cannot. An example of a potent potentiator of glucose-induced insulin secretion is 2-BFI. However, it requires higher concentrations than what is typical for interactions with I₂Rs. Other ligands of I₂Rs, such as idazoxan and RS45041-190, do not induce insulin release or antagonize the insulin secretory effect of other imidazoline compounds on b-cells.^{36,37} It seems that the presence of 2-BFI and its related compounds does not have a direct effect on b-cell insulin secretion through I₂Rs. One possible

explanation for how imidazoline compounds alter insulin secretion is that they bind to a site that is linked to the ATP-sensitive potassium (KATP) channel, which then lowers the rate of potassium efflux. This, in turn, increases the membrane potential and leads to membrane depolarization, ultimately triggering insulin secretion.³⁴ The b-cell κ ATP channel is made up of two subunits, Kir6.2 and sulfonylurea receptor 1 (SUR1). Previous research has demonstrated that imidazolines can still function as blockers of KATP channels even in the absence of SUR1, when truncated Kir6.2 is present. This suggests that SUR1 is not necessary for imidazoline compounds to regulate the ionic conductance of Kir6.2. Additionally, the study indicates that the I₃Rs are situated within Kir6.2.^{38,39} Further biochemical evidence supports this view.⁴⁰ Therefore, I₃Rs represent a binding site located on the Kir6.2 subunit of κ ATP channels. I₃Rs were found in the pancreas by sensitivity to the imidazoline receptor agonist, efaroxan. Also, one of the imidazoline derivatives, diakamph (V), showed clear hypoglycemic activity and was positioned as a treatment for type 2 diabetes mellitus. In addition, it was found to have neuroprotective effects with oxidative cytotoxicity. Cytoprotective activity against cerebral ischemia in rats has been shown for other IR agonists.⁴¹ Agmatine, considered a natural agonist of IRb in experiment, showed antidepressant activity and exhibited anxiolytic, antinociceptive and anti-inflammatory properties.^{8,42-44}

Among the identified receptors/enzymes for imidazoline derivatives, the following have been reliably confirmed: for I₂R - monoamine oxidases A, B, and brain b-creatinine kinase,⁴⁵ for I₁R - nischarin.⁴⁶ Another benzimidazole derivative – bendazole inhibits experimental myopia progression and decreases the ocular accumulation of HIF-1 α protein in young rabbits.⁴⁷

Bendazole⁴⁷ and diakamph⁴⁸ are not pure imidazolines but only contain its fragment in the form of benzimidazole, while they exhibit more universal effects (hypotensive, sugar-reducing, adaptogenic, neuroprotective) all 3 types of imidazoline receptors, which gave us grounds to choose this group of compounds for modeling, synthesis and study of biological activity in relation to models of pathologies, in the treatment of which imidazoline receptor agonists have been successfully used.

Benzimidazole derivatives are crucial in many biologically active compounds, including antiparasitics, antimicrobials, antivirals, antifungals, and more. Drugs in various therapeutic lines have been developed by changing substituents around the core structure, such as albendazole, carbendazim, and omeprazole.⁴⁹

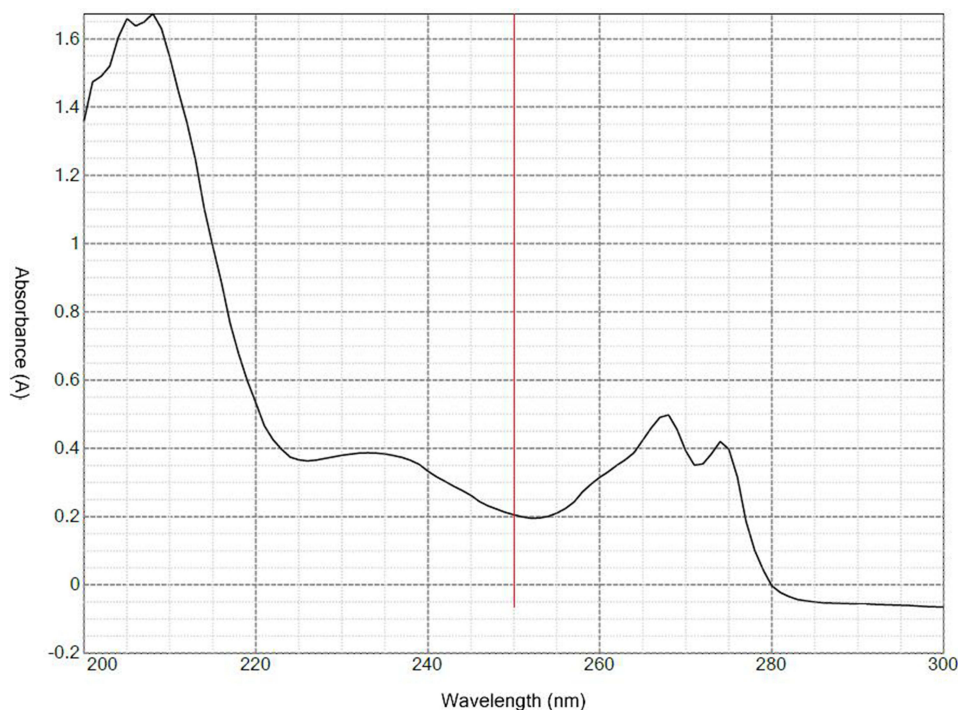


Figure 2 UV-specter of (IVd) benzimidazole derivative.

The improved methodology for synthesizing benzimidazoles developed by us made it possible to obtain derivatives of thermolabile carboxylic acids, including those from amino acids and their derivatives, such as lipoic acid, in a more gentle regime.

The essence of the technique consists in boiling the reagents in glacial acetic acid with a reflux condenser and a Dean-Stark nozzle. After distillation of 2/3 of glacial acetic acid, the Dean-Stark nozzle is removed, and the reaction mixture continues to be heated for another 3–10 hours. This technique allows the use of hydrophilic acids and salts of amino acids as donors of carboxyl groups. Older methods involve using hydrophobic solvents such as benzene and toluene at higher boiling points. In this case, the product yield is minimal (up to 15%), some initial acids are decarboxylated, and some are simply not dissolved in toluene and benzene. This did not allow previously to obtain benzimidazoles from thermolabile or bifunctional compounds.

Materials and Methods

Experimental Section

The molecular docking method predicts how a protein binds to a ligand. Docking algorithms determine the best orientation with the lowest energy score.⁵⁰

Computational Analysis

Ligand and Protein Preparation Ligands

As ligands for modeling, we used: benzimidazolyl-arginine (IVa), benzimidazolyl-lysine (IVb), benzimidazolyl-threonine (IVc), benzimidazolyl-histidine (IVd), benzimidazolyl-thioctic acid (IVe), benzimidazolyl-ursodeoxycholic acid (IVf), diakamph, moxonidine). As controls with known ability to stimulate all three types of imidazoline receptors, we used diakamph and moxonidine.

Molecular Docking

Diakamph (V) and moxonidine (VI) were obtained from PubChem database,⁵¹ the other substances were constructed using Yasara program,⁵² and their geometric optimization was performed using AMBER 15 IPQ method program,⁵³ which includes geometric optimization in a water box with alternating sessions of molecular dynamics to select the most stable conformer with minimum energy. Protein obtained from PDB database (<https://www.rcsb.org/>): model monoamine oxidase-B (2BK3) in complex with the agonist farnesol, monoamine oxidase B in complex with the agonist (2-benzofuranyl)-2-imidazoline (2XCG), chicken brain creatinine kinase (1QH4) and human discharging PX-domain (3P0C).

Yasara software (<http://www.yasara.org/>) was used for docking after removing all water and ligand molecules from the protein preparation.

Previously, a comparative study of molecular docking methods was performed and showed that the methods' results are comparable to each other and give statistically comparable results. Multiligand docking and the highest output parameters are given by the iGemdock program (<http://gemdock.life.nctu.edu.tw/dock/igemdock.php>). In addition, this program also takes pharmaceutical semi-empirical criteria such as PharmScore into account.

Molecular docking software used in iGemdock 2.1. iGemdock application, first, we have to select the protein binding site and ligands; virtual screening procedure of iGemdock consists of four main steps (Population size = 200, Generations = 70, Number of solutions = 2, and Default setting = Standard docking).^{54,55} iGemdock suite is an automated docking/screening tool used a generic evolutionary method for molecular docking and empirical scoring function. iGemdock useful tool for molecular recognition and used to systematically evaluate and improve docking scoring function.^{56,57} We used the PyMOL program (<https://pymol.org/2/>) to create a 3D model of the active centers with the ligands.^{58,59} Additionally, we utilized the LigPlus program (<https://www.ebi.ac.uk/thornton-srv/software/LigPlus/>) to generate 2D projections of the complexes.^{60,61}

Materials and Equipment

All chemicals and reagents used in the study were obtained from commercial vendors, including Merck (Germany) and Sigma Aldrich Sigma Chemical, Co. (USA), and were utilized without further purification. HPLC was performed using an HPLC microcolumn chromatograph (Millichrom A-02, RF)^{62,63} with a gradient of acetonitrile (5–100%)/0.1 M chloric acid and 0.5 M lithium perchlorate).

Melting points were determined using the open capillary method in an uncorrected Gallen Kamp MFB-595^{64,65} electric melting point instrument. UV spectra were obtained using a double-beam UV-Visible spectrophotometer GeneQuant 1300 (Model 2021, UK).^{66,67}

The ¹H-NMR, ¹³C-, NMR spectra were obtained using a Bruker Avance III 400^{68,69} at 400 and 100 MHz, with Tetramethylsilane (TMS) as an internal standard. The spectra were measured relative to TMS in DMSO-d₆. The chemical shifts are represented as s (singlet), d (doublet), t (triplet), q (quartet), m (multiplet), dd (doublet of doublets), and br s (broad singlet) in the appropriate solvent, along with coupling constants (J) in Hz (Hertz).

(R)-1-(4-Amino-4-(1H-Benzo[D]imidazol-2-Yl)butyl)guanidine (IVa)

L-arginine (base) (174 mg, 1.0 mmol) and benzene-1,2-diamine (108 mg, 1.0 mmol), dissolve in 3 mL anhydrous glacial acetic acid (AGAA) in a 10 mL round bottom flask, place in a system with a reflux condenser and a Dean-Stark nozzle. Boil with the reflux condenser and extract 2 mL of AGAA. Then, the Dean-Stark nozzle is disconnected, and the solution is boiled with reflux condenser for 5 hours. The solution is cooled, 6 mL of ethanol is added and placed in a -15 ° C freezer. The precipitate (IVa) is filtered off and recrystallized from a 1:3 mixture of AGAA and ethanol, and the precipitate is dried. (S)-1-(4-(4-amino-4-(1H-benzo[d]imidazol-2-yl)butyl) guanidine (IVa) is obtained as a gray powder. For analytical purposes, the product (IVa) is additionally was purified by prep-HPLC (Phenomenex Luna C18 100 mm × 40 mm × 5 μm,⁷⁰ mobile phase: [water (0.1%TCIA) – ACN]: 20–55%, 12 min). ¹H-NMR (400 MHz, DMSO-d₆): δ 1.50 (m, 2H), 1.81 (m, 2H), 2.52 (s, 1H), 3.33 (m, 2H), 3.8 (s, 1H), 7.16 (d, 1H), 7.52 (s, 1H), 8.76 (s, 2H), 6.63 (s, 2H), 7.82 (s, 1H), 12.16 (s, 1H). ¹³CNMR (126 MHz, CD₃OD) δ 24.39, 25.70, 30.66, 43.07, 58.05, 58.66, 79.19, 100.37. HRMS (ESI +): m/z calcd for C₁₂H₁₈N₆: 246.3212 [M + H]⁺; found: 246.33188 [M + H]⁺.

(S)-1-(1H-Benzo[D]imidazol-2-Yl)pentane-1,5-Diamine (IVb)

L-lysine (base) (146 mg, 1.0 mmol) and benzene-1,2-diamine (108 mg, 1.0 mmol), dissolve in 5 mL anhydrous glacial acetic acid (AGAA) in a 20 mL round bottom flask, place in a system with a reflux condenser and a Dean-Stark nozzle. Boil with the reflux condenser and extract 3 mL of AGAA. Then, the Dean-Stark nozzle is disconnected, and the solution is boiled with reflux condenser for 5 hours. The solution is cooled, 9 mL of ethanol is added and placed in a -15 ° C freezer. The precipitate (IVb) is filtered off and recrystallized from a 1:3 mixture of AGAA and ethanol, and the precipitate is dried. (S)-1-(1H-benzo[d]imidazol-2-yl)pentane-1,5-diamine (IVb) is obtained as a gray-yellow powder. For analytical purposes, the product (IVb) is additionally was purified by prep-HPLC (Phenomenex Luna C18 100 mm × 40 mm × 5 μm, mobile phase: [water (0.1%TCIA) – ACN]: 20–55%, 12 min). ¹H-NMR (400 MHz, DMSO-d₆): δ 1.19 (m, 2H), 1.52 (m, 2H), 1.81 (m, 2H), 1.91 (m, 2H), 2.74 (dd, 2H), 3.95 (br s, 1H), 4.34 (m, 1H), 7.32 (2dd, 1H). ¹³CNMR (126 MHz, CD₃OD) δ 22.09, 24.39, 27.69, 30.66, 41.90, 58.06, 58.66, 79.20, HRMS (ESI+): m/z calcd for C₁₂H₁₈N₄: 218.3011 [M + H]⁺; found: 218.31149 [M + H]⁺.

(1R,2R)-1-Amino-1-(1H-Benzo[D]imidazol-2-Yl)propan-2-Ol (IVc)

L-threonine (119 mg, 1.0 mmol) and benzene-1,2-diamine (108 mg, 1.0 mmol), dissolve in 3 mL anhydrous glacial acetic acid (AGAA) in a 10 mL round bottom flask, place in a system with a reflux condenser and a Dean-Stark nozzle. Boil with the reflux condenser and extract 2 mL of AGAA. Then, the Dean-Stark nozzle is disconnected, and the solution is boiled with reflux condenser for 5 hours. The solution is cooled, 6 mL of methanol is added and placed in a -15 ° C freezer. The precipitate (IVc) is filtered off and recrystallized from a 1:3 mixture of AGAA and methanol, the precipitate is dried (1R,2R)-1-amino-1-(1H-benzo[d]imidazol-2-yl)propan-2-ol (IVc) is obtained as a light gray powder. For analytical purposes, the product (IVc) is additionally was purified by prep-HPLC (Phenomenex Luna C18 100 mm × 40 mm × 5 μm, mobile phase: [water (0.1%TCIA) – ACN]: 20–55%, 4 min). ¹H-NMR (400 MHz, DMSO-d₆): δ 1.11 (m, 3H), 3.60 (br s, 2H), 3.81 (d, 1H), 4.00 (m, 1H), 7.30 (m, 1H), 7.35 (m, 1H), 3.95 (br s, 1H). ¹³CNMR (126 MHz, CD₃OD) δ 20.19, 24.39, 30.66, 41.90, 58.05, 58.66, 67.75, 79.20, HRMS (ESI+): m/z calcd for C₁₀H₁₃N₃O: 191.2302 [M + H]⁺; found: 191.2050 [M + H]⁺.

1-(1H-Benzo[D]imidazol-2-Yl)-2-(1H-Imidazol-4-Yl)ethan-1-Amine (IVd)

Histidine (155 mg, 1.0 mmol) and benzene-1,2-diamine (108 mg, 1.0 mmol), dissolve in 5 mL anhydrous glacial acetic acid (AGAA) in a 20 mL round bottom flask, place in a system with a reflux condenser and a Dean-Stark nozzle. Boil with the reflux condenser and extract 3 mL of AGAA. Then, the Dean-Stark nozzle is disconnected, and the solution is boiled with reflux condenser for 7 hours. The solution is cooled, 6 mL of ethanol is added and placed in a -15°C freezer. The precipitate (IVd) is filtered off and recrystallized from a 1:3 mixture of AGAA and methanol, the precipitate is dried 1-(1H-benzo[d]imidazol-2-yl)-2-(1H-imidazol-4-yl)ethan-1-amine (IVd) is obtained as a blue powder. For analytical purposes, the product (IVd) is additionally was purified by prep-HPLC (Phenomenex Luna C18 100 mm \times 40 mm \times 5 μm , mobile phase: [water (0.1%TCIA) – ACN]: 20–55%, 4 min). 1H-NMR (400 MHz, DMSO-d₆): δ 2.61 (m, 2H), 2.86 (dd, 2H), 2.90 (d, 1H), 4.75 (d, 1H), 4.77 (d, 1H), 5.24 (m, 1H), 7.33 (m, 1H). 13CNMR (126 MHz, CD₃OD) δ 29.21, 58.05, 114.80, 115.90, 117.44, 122.45, 131.74, 135.74, 137.29, 152.89. HRMS (ESI+): m/z calcd for C₁₀H₁₃N₃O: 227.2702 [M + H]⁺; found: 227.2212 [M + H]⁺.

(R)-2-(4-(1,2-Dithiolan-3-Yl)butyl)-1H-Benzo[D]imidazole (IVe)

Lipoic acid ((R)-5-(1,2-dithiolan-3-yl)pentanoic acid) (206 mg, 1.0 mmol) and benzene-1,2-diamine (108 mg, 1.0 mmol), dissolve in 5 mL anhydrous glacial acetic acid (AGAA) in a 20 mL round bottom flask, place in a system with a reflux condenser and a Dean-Stark nozzle. Boil with the reflux condenser and extract 3 mL of AGAA. Then, the Dean-Stark nozzle is disconnected, and the solution is boiled with a reflux condenser for 5 hours. The solution is cooled, 4 mL of anhydrous dioxan is added and placed in a -15°C freezer. The precipitate (IVe) is filtered off and recrystallized from a 1:2 mixture of AGAA and dioxan, and the precipitate is dried (R)-2-(4-(1,2-dithiolan-3-yl)butyl)-1H-benzo[d]imidazole (IVe) is obtained as a yellow powder. For analytical purposes, the product (IVe) is additionally was purified by prep-HPLC (Phenomenex Luna C18 100 mm \times 40 mm \times 5 μm , mobile phase: [water (0.1%TCIA) – ACN]: 20–55%, 17 min). 1H-NMR (400 MHz, DMSO-d₆): δ 1.34 (d, 2H), 1.36 (d, 2H), 1.38 (s, 2H), 1.41 (s, 1H), 1.48 (m, 2H), 1.70 (d, 2H), 1.72 (d, 2H), 2.50 (m, 2H), 2.78 (s, 2H), 2.78 (s, 2H), 2.81 (d, 2H), 3.70 (m, 1H), 3.70 (m, 1H), 7.00 (m, 1H), 7.01 (m, 1H), 7.35 (m, 1H). 13CNMR (126 MHz, CD₃OD) δ 26.03, 27.19, 27.73, 34.31, 38.33, 40.13, 56.21, 114.81, 115.91, 122.46, 122.54, 138.00, 155.71. HRMS (ESI+): m/z calcd for C₁₄H₁₈N₂S₂: 278.4305 [M + H]⁺; found: 278.4211 [M + H]⁺.

(3R,5S,7S,9S,10S,13R,14S,17R)-17-((R)-4-(1H-Benzo[D]imidazol-2-Yl)butan-2-Yl)-10,13-Dimethylhexadecahydro-1H-Cyclopenta[a]phenanthrene-3,7-Diol (IVf)

Ursodesoxycholic acid ((4R)-4-((3R,5S,7S,9S,10S,13R,14S,17R)-3,7-dihydroxy-10,13-dimethylhexadecahydro-1H-cyclopenta[a]phenanthren-17-yl)pentanoic acid) (392 mg, 1.0 mmol) and benzene-1,2-diamine (108 mg, 1.0 mmol), dissolve in 30 mL anhydrous glacial acetic acid (AGAA) in a 50 mL round bottom flask, place in a system with a reflux condenser and a Dean-Stark nozzle. Boil with the reflux condenser and extract 20 mL of AGAA. Then, the Dean-Stark nozzle is disconnected, and the solution is boiled with reflux condenser for 10 hours. The solution is cooled, 4 mL of methanol is added and placed in a -15°C freezer. The precipitate (IVf) is filtered off and recrystallized from a 1:3 mixture of AGAA and methanol, the precipitate is dried (3R,5S,7S,9S,10S,13R,14S,17R)-17-((R)-4-(1H-benzo[d]imidazol-2-yl)butan-2-yl)-10,13-dimethylhexadecahydro-1H-cyclopenta[a]phenanthrene-3,7-diol (IVf) is obtained as a white, yellow powder. For analytical purposes, the product (IVf) is additionally was purified by prep-HPLC (Phenomenex Luna C18 100 mm \times 40 mm \times 5 μm , mobile phase: [water (0.1%TCIA) – ACN]: 20–55%, 17 min). 1H-NMR (400 MHz, DMSO-d₆): δ 0.70 (s, 3H), 0.95 (s, 3H), 1.32 (s, 2H), 1.47 (m, 2H), 1.58 (m, 2H), 1.61 (m, 1H), 1.81 (m, 2H), 2.03 (m, 2H), 2.73 (d, 2H), 2.75 (d, 2H), 3.65 (br s, 1H), 3.74 (br s, 1H), 4.38 (br s, 1H), 7.02 (m, 1H), 7.34 (m, 1H). 13CNMR (126 MHz, CD₃OD) δ 12.75, 18.89, 20.00, 22.40, 24.39, 24.89, 28.32, 28.84, 30.52, 30.66, 31.26, 34.19, 35.69, 35.83, 36.06, 36.39, 40.33, 40.40, 40.80, 42.63, 43.89, 51.80, 56.87, 58.66, 68.84, 71.30, 79.20. HRMS (ESI+): m/z calcd for C₃₀H₄₄N₂O₂: 464.3412 [M + H]⁺; found: 464.3600 [M + H]⁺.

The data are presented in Table 5.

Results and Discussion

During the synthesis of each derivative (IVa-f), certain peculiarities were observed. The highly hydrophilic substances (IVa-d) caused the solution to turn green and dark blue during synthesis. No precipitates formed in the solutions after stopping the reaction and cooling the mixture. As a result, it was necessary to add ethanol or isopropanol and sometimes freeze the solution to obtain the residue. Recrystallization of substances (IIa-d) also involved adding ethanol or isopropanol to a solution of the derivative in glacial acetic acid, followed by freezing the solution.

After adding substances (IVe,f), the reaction mixture turned light blue and formed blue crystals upon settling. Based on the liquid chromatography data, the reaction time spanned from 2 to 8 hours, with peak 4 having the maximum area and peaks 1 and 3 decreasing in area. The figure displays the chromatogram of the reaction mixture during the product (IVd) after the solution acquired coloration. Band 1 is histidine, band 2 is probably the salts of o-phenylenediamine (I), histidine, and acetic acid, band 3 corresponds to the amide of histidine and o-phenylenediamine, and band 4 indicates the final product (IVd).

Preparative chromatographic purification gives one absorption band corresponding to the UV spectrum of benzimidazole with a split peak in the region 260–280 nm (Figure 2). A similar situation is observed in the synthesis of the derivative (IIc): bands 1 and 2 are mixtures of the starting components serine (IIc) and o-phenylenediamine (I), band two is associated with amides (IIIc), and band 4 is the final product (IVc).

The synthesis of diakamph, which was described in,⁷⁹ was used as an example for comparison. Figures 4–6 show the results of RP-HPLC analysis of the reaction mixture. The mixture contains (I) and amides associated with band 2, as well as a mixture of diakamph isomers with bands 3 and 4. The spectral ratios for bands 4 in all the chromatograms exhibit a double.

Table 1 Dynamics of Blood Glucose in Rats with Alloxan Diabetes After a Single Oral Administration of an Benzimidazoles IVa-f

Glucose Content in Blood Serum. mmol / l						
Groups of Animals	Substance	Initial Level	1 h	2 h	3 h	24 h
1	–	4.22±0.24	4.52±0.47	4.40±0.42	4.36±0.47	4.40±0.45
2	IVa	4.28±0.34	4.90±0.51	4.30±0.26	4.24±0.47	4.58±0.38
3	IVb	4.68±0.65	4.60±0.27	4.42±0.28	4.40±0.29	4.58±0.51
4	IVc	4.34±0.58	4.62±0.40	4.54±0.55	4.54±0.40	4.40±0.51
5	IVd	4.68±0.29	4.66±0.34	4.48±0.40	4.94±0.15	4.36±0.53
6	IVe	4.84±0.68	4.52±0.58	4.70±0.30	4.70±0.25	4.72±0.41
7	IVf	4.40±0.52	4.40±0.33	4.50±0.42	4.52±0.41	4.58±0.41
8	–	16.72±1.88	17.06±1.38	17.16±1.36	17.58±1.85	18.12±1.80
9	IVa	12.36±0.89	11.06±0.19*	8.94±0.50*	8.28±1.11*	8.30±0.39*
10	IVb	16.02±1.76	15.94±1.62	15.70±1.60	14.78±1.36	16.74±1.90
11	IVc	14.36±2.36	14.66±1.04	15.06±1.52	16.28±2.69	16.06±1.53
12	IVd	11.94±0.63	10.84±0.66*	9.84±0.36*	9.10±0.62*	8.54±0.76*
13	IVe	12.56±1.43	14.20±1.30	15.18±0.90	16.60±0.77	17.70±1.33
14	IVf	12.22±0.85	9.14±2.03*	8.12±0.93*	8.16±0.53*	7.82±0.42*

Notes: *Differences are statistically reliable between this group and the control group N 8 (P≤0.05); there are no statistical differences between groups N 1–7; control group N 8 with diabetes is statistically significantly different from groups 1–7 (P≤0.05).

Table 1 (Full data is shown in [Supplementary Material](#)) shows that giving derivatives (IVa-f) to animals did not affect their blood glucose levels. The average glucose level in groups 2–7 was the same as the control group (N 1). In the control group of animals with induced diabetes (N 8), glucose levels were between 16.7 and 18.1 mmol/l, which is typical for diabetes mellitus. A significant reduction in blood sugar levels ($P \leq 0.05$) was observed for compounds (IVa), (IVd), and (IVf). Among these three compounds, the benzimidazole derivative and ursodeoxycholic acid (IVf) had the highest sugar-reducing effect and the lowest energy in docking.

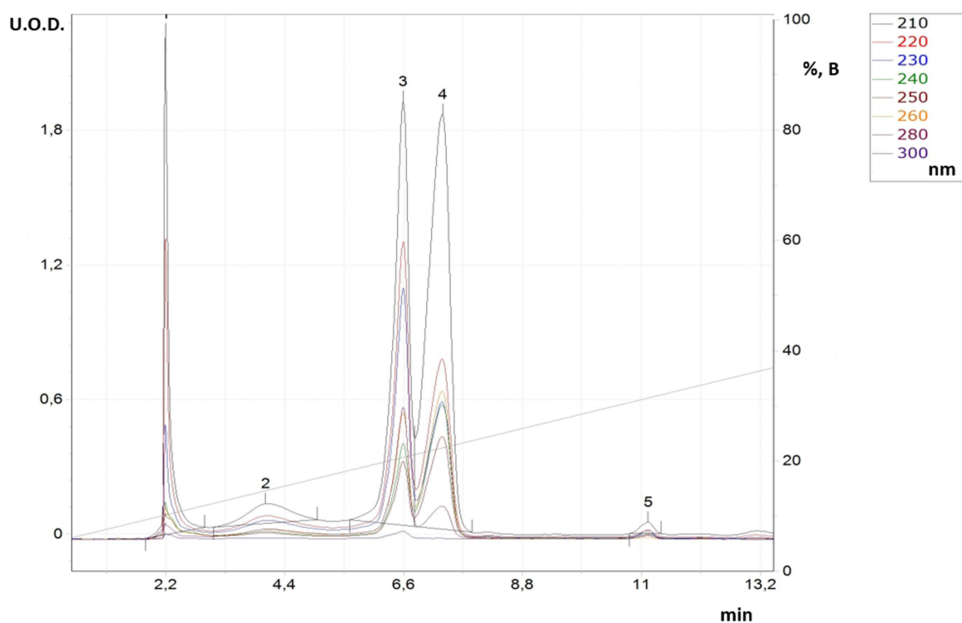


Figure 4 RP-HPLC for reaction solution in synthesis process for derivative (IVd).

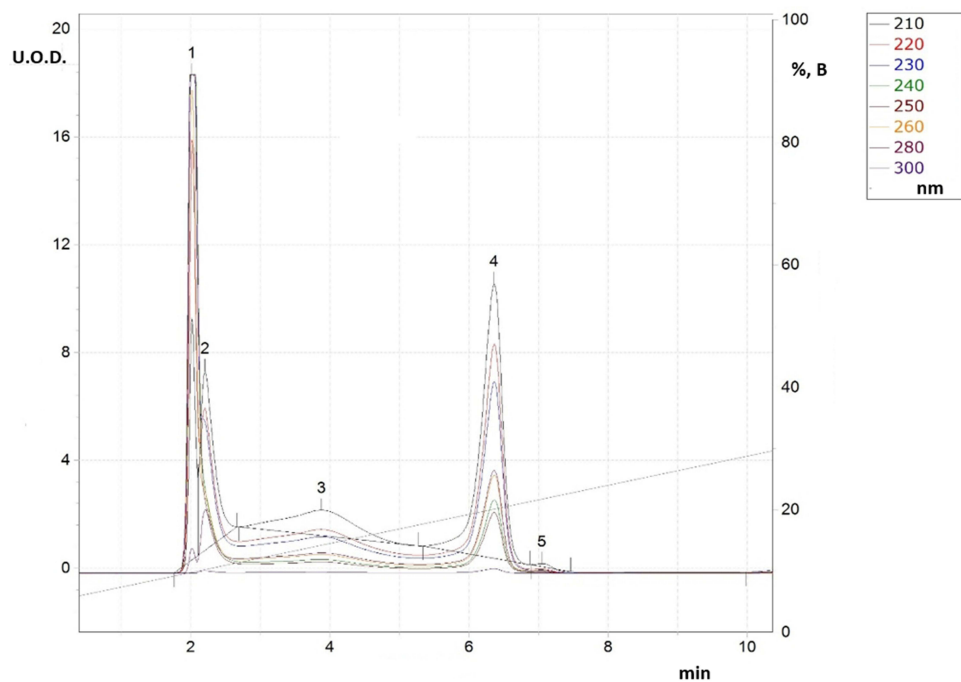


Figure 5 RP-HPLC for reaction solution in synthesis process for derivative (IVc).

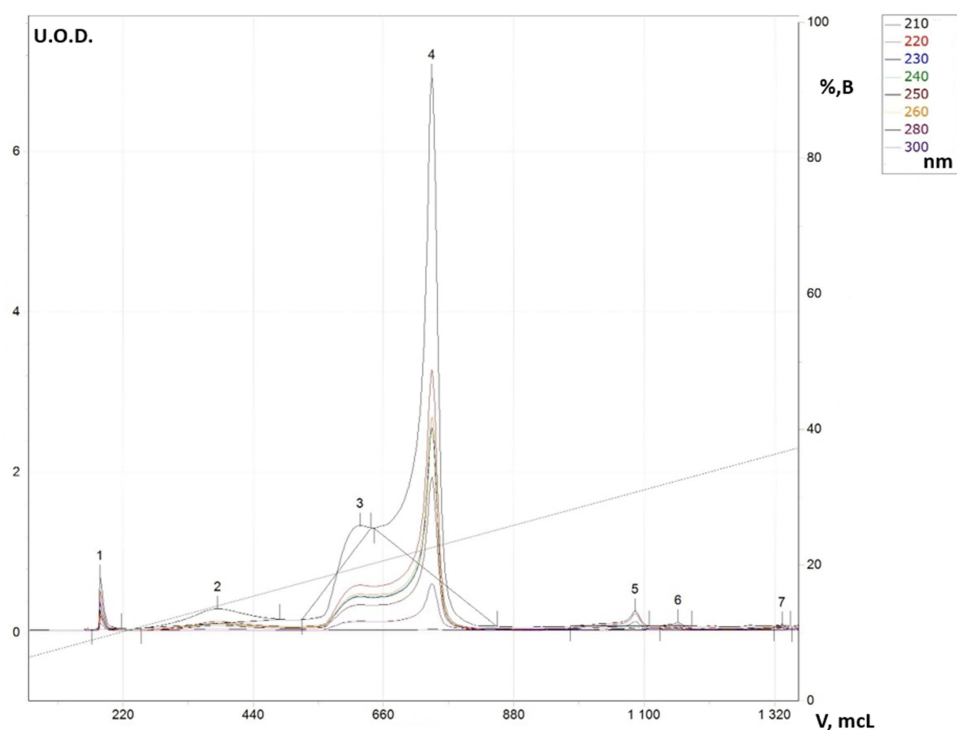


Figure 6 RP-HPLC for reaction solution in synthesis process for diacmph (V).

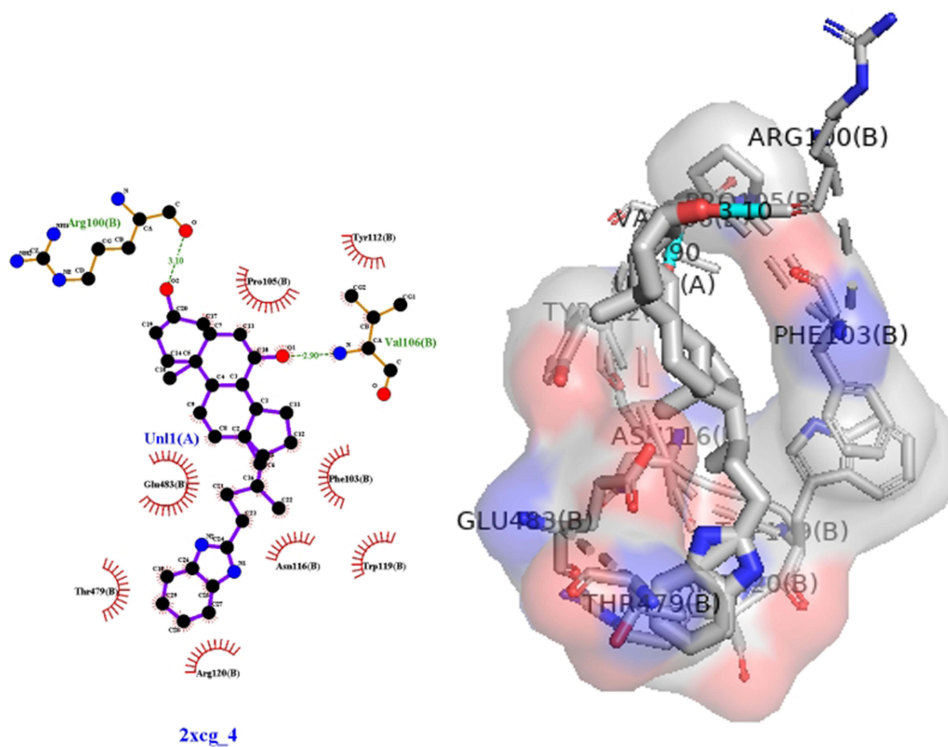


Figure 7 3D (right) and 2D (left) representations of the binding interactions of (IVf) against Human monoamine oxidase b (PDB ID: 2XCG).

Table 2 Results of Molecular Docking Between Studied Substances and Chicken Brain-Type Creatine Kinase (PBP) (PDB ID: 1QH4)

Ligand	Total Energy, kcal/mol	VDW ^a	HBond ^b	Elec ^c	AverConPair ^d
IVa	-87.9522	-63.4528	-24.5208	0.0214502	24.3889
V	-81.812	-65.199	-16.613	0.00	36.00
VI	-76.0144	-63.9707	-12.0437	0.00	27.6667
IVd	-78.0442	-63.0938	-14.9504	0.00	27.3125
IVe	-79.8191	-66.0259	-13.7932	0.00	30.75
IVb	-71.1517	-50.2328	-20.9189	0.00	25.6429
IVc	-81.812	-65.199	-6.79144	0.00	19.8235
IVf	-101.363	-94.572	-16.613	0.00	36.00

Notes: ^aVan der Waals Potential in Protein Complexes; ^bSolvation energy of hydrogen bond; ^cElectrostatic potential; ^dAverage confirmation pair.

The synthesized compounds (IVa-f) have minimum binding energy (Table 2) ranging from (-70 to -107 kcal/mol) with a good result achieved using compound 4 (-106.6 kcal/mol), which revealed Figure 3 3D (right) and 2D (left) representations of the binding interactions of (IVf) against Human monoamine oxidase b (PBP) (PDB ID: 2XCG). Compound (IVf) interacts Arg-100 was involved in hydrogen bonding with -C=O of (IVf) and with NH₂-group of Val-106. Hydrophobic/ π -cation interactions were observed for Tyr-112 with the aromatic group, while Arg-120, Thr-479, Glu-483, Asn-116, Trp-119, Tyr-112 and Pro-105 showed Van der Waals residual interactions, respectively (Figure 7). The lowest energy of target-ligand complexes, except for compound (IVf), was observed in compounds (IVa) and the control substance II-stimulant agonist moxonidine (VI). This suggests the presence of sugar-lowering activity in the synthesized substances.

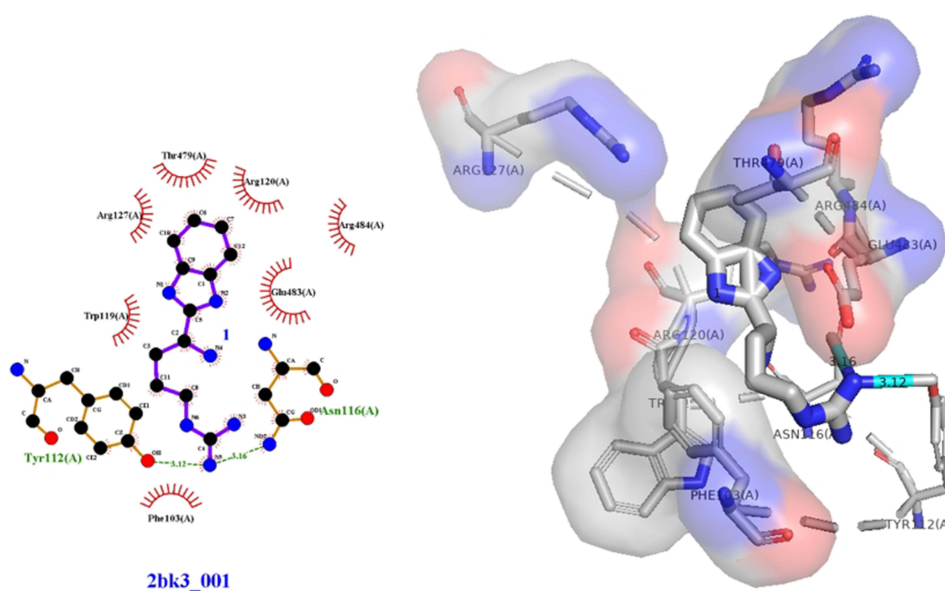
**Figure 8** 3D (right) and 2D (left) representations of the binding interactions of (IVa) against Human monoamine oxidase b (PBP) (PDB ID: 2bk3).

Table 3 Results of Molecular Docking Between Studied Substances and Human Monoamine Oxidase b (PBPs) (PDB ID: 2XCG)

Ligand	Total Energy, kcal/mol	VDW ^a	HBond ^b	Elec ^c	AverConPair ^d
IVa	-106.588	-85.7413	-21.4857	0.638991	33.6667
V	-83.335	-58.0035	-25.3315	0	27.0500
VI	-99.6059	-83.6347	-13.6195	-2.35168	32.75
IVd	-90.6571	-79.9717	-10.6853	0	33.9333
IVe	-91.1353	-89.5565	-1.57887	0	31.8333
IVb	-94.3503	-84.0812	-10.2691	0	34.0000
IVc	-87.5584	-72.3225	-15.2359	0	32.7857
IVf	-103.305	-92.2593	-11.0457	0	30.0882

Notes: ^aVan der Waals Potential in Protein Complexes; ^bSolvation energy of hydrogen bond; ^cElectrostatic potential; ^dAverage confirmation pair.

Compound (IVa) interacts Asn-116 was involved in hydrogen bonding with -NH₂ of (IVa) and with OH-group of Tyr112. Hydrophobic/ π -cation interactions were observed for Phe-103 with the aromatic group, while Trp-119, Arg-127, Thr-479, Arg-120, Arg-484 and Glu-483 showed Van der Waals residual interactions, respectively (Figure 8). Among other molecules, (IVa) had one of the lowest complex energies with the 2bk3 target -94.5 kJ/mol (Table 3).

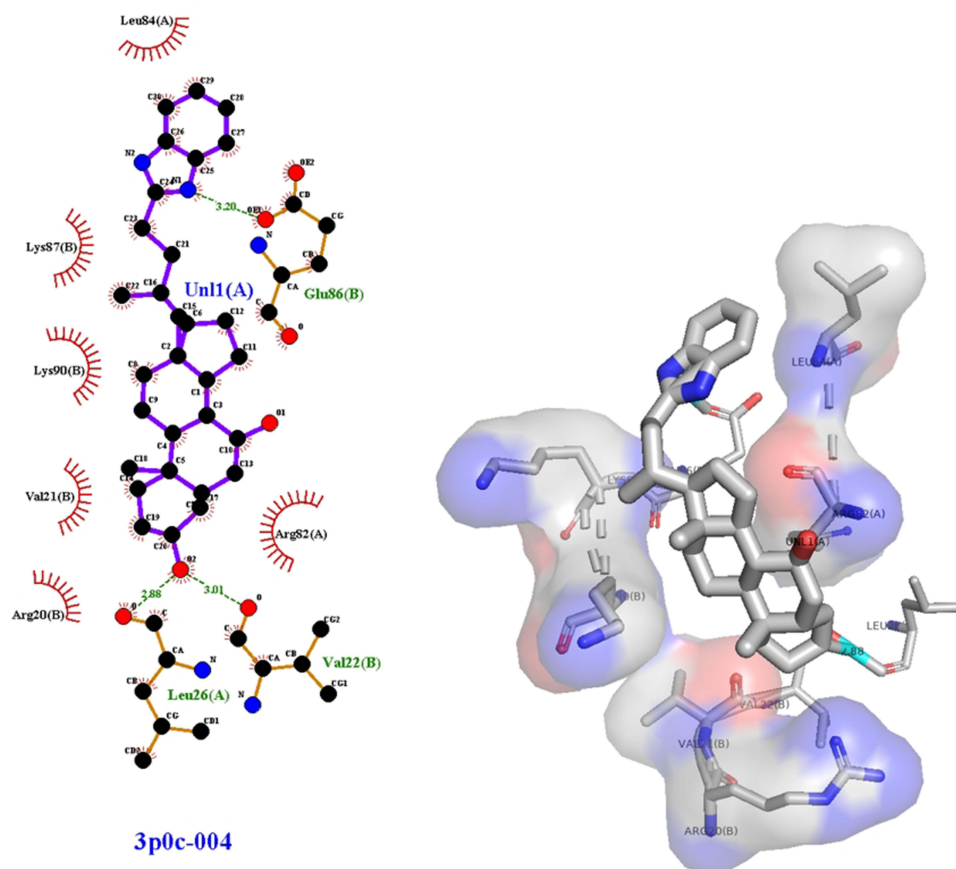
**Figure 9** 3D (right) and 2D (left) representations of the binding interactions of (IVf) against nischarin (PBPs) (PDB ID: 3p0c).

Table 4 Results of Molecular Docking Between Studied Substances and Human Monoamine Oxidase b (PBP) (PDB ID: 2bk3)

Ligand	Total Energy, kcal/mol	VDW ^a	HBond ^b	Elec ^c	AverConPair ^d
IVa	-94.4941	-68.2404	-25.4673	-0.786414	29.3889
V	-96.7332	-82.4078	-14.3254	0	45.165
VI	-89.1305	-72.3257	-16.8048	0	31.7333
IVd	-93.7606	-85.8706	-7.88996	0	30.8333
IVe	-91.2754	-71.4504	-19.825	0	28.0625
IVb	-85.3885	-79.1634	-6.22513	0	31.25
IVc	-88.3253	-73.4629	-14.8624	0	33.2857
IVf	-102.513	-101.436	-1.07632	0	21.5882

Notes: ^aVan der Waals Potential in Protein Complexes; ^bSolvation energy of hydrogen bond; ^cElectrostatic potential; ^dAverage confirmation pair.

Compound (IVf) interacts with Leu-26 through ionic bonds -CH₂-OH of (IVf) and with -C=O-group of Val-22. An ionic bond is also formed between the -COO group in Glu-86 and the nitrogen of (IVf) imidazole fragment. Leu-84, Lys-87, Lys-90, Val-21, Arg-20 and Arg-82 showed Van der Waals residual interactions, respectively (Figure 9). The energy of the complex of Human Nischarin (PBP) (PDB ID: 3P0C) with the derivative (IVf) was minimal -101.4 kJ/mol (Table 4). Of all compounds, the most promising 3P0C agonists can be considered compounds (IVa) and the comparison substance diakamph (V) with proven antagonism to nischarin.

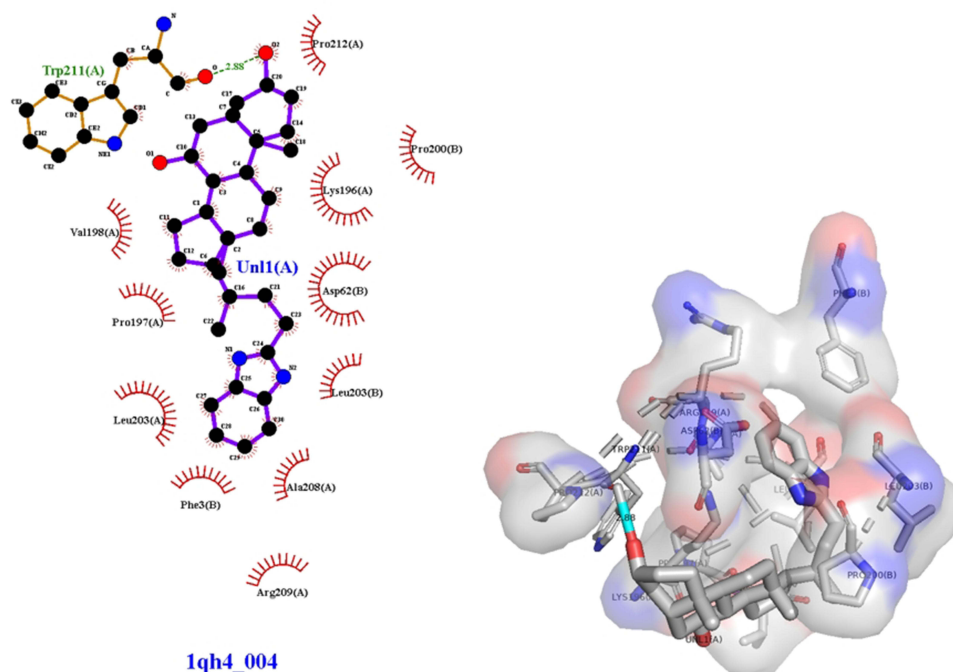
**Figure 10** 3D (right) and 2D (left) representations of the binding interactions of (IVf) against chicken brain-type creatine kinase (PBP) (PDB ID: 1QH4).

Table 5 Results of Molecular Docking Between Studied Substances and Nischarin (PBPs) (PDB ID: 3p0c)

Ligand	Total Energy, kcal/mol	VDW ^a	HBond ^b	Elec ^c	AverConPair ^d
IVa	-82.2266	-55.4693	-26.7573	0	22.5556
V	-67.3746	-53.4233	-13.9513	0	22
VI	-66.5056	-56.0135	-10.492	0	18.5556
IVd	-70.8207	-56.8219	-13.9988	0	23.9375
IVe	-67.9041	-51.7674	-16.1367	0	22
IVb	-68.8429	-47.3943	-21.4486	0	21.5714
IVc	-82.804	-69.816	-12.988	0	13.7353
IVf	-67.3746	-53.4233	-13.9513	0	22

Notes: ^aVan der Waals Potential in Protein Complexes; ^bSolvation energy of hydrogen bond; ^cElectrostatic potential; ^dAverage confirmation pair.

Figure 10 3D (right) and 2D (left) representations of the binding interactions of (IVf) against chicken brain-type creatine kinase (PBPs) (PDB ID: 1QH4). Compound (IVf) interacts with Trp-211 through ionic bonds -C=O and -OH-02 in (IVf). Val-198, Pro-197, Lys-90, Leu-203, Phe-3, Arg-209, Ala-208, Leu-203, Asp-62, Lys-196, Pro-200, Pro-212 showed Van der Waals residual interactions, respectively (Figure 10). The energy of the complex of chicken brain-type creatine kinase (PBPs) (PDB ID: 1QH4) with the best pose-derivative (IVf) was minimal -101.4 kJ/mol (Table 5). Of all compounds, the most promising 1QH4 agonists can be considered compounds (IVa) and the comparison substance diakamph (V) with proven antagonism to nischarin.

Conclusions

1. Benzimidazole derivatives based on thermolabile amino acids (arginine, lysine, serine, histidine) and natural organic acids (ursodeoxycholic and lipoic acids) were synthesized.
2. Molecular modeling was used to predict the highest affinity to imidazoline receptors for derivatives (IVa) and (IVf). Among these, the derivative (IVf) based on ursodeoxycholic acid displayed the lowest energies of complexes with imidazoline receptors of both type 1 and types 2 and 3. This derivative has the potential to be developed as a new antihyperglycemic agent for the treatment of severe forms of diabetes.
3. The synthesized compounds were studied on the model of alloxan diabetes in rats. As a result, compounds (IVa), (IVd), and (IVf) showed a significant and statistically proven antihyperglycemic effect. Among these compounds, the ursodeoxycholic acid derivative (IVf) had the most pronounced effect, as predicted by the docking method.
4. In the future, modeling is planned to provide molecular dynamic simulation data and thermodynamic stabilities of Ligand-Protein complex for the complex IVf-protein and Pharmacokinetic and bioavailability predictions.
5. It is also planned to study the toxic properties of the most promising compounds (IVa) and (IVf).

Institutional Review Board Statement

The animals were cared for according to the international guidelines 33647-2015 (Principles of the Good Laboratory Practice, GLP), the international recommendations of the European Convention for the Protection of Vertebrate Animals used for Experimental and Other Scientific Purposes (The European Convention, 1986). The protocol of the experimental study was approved by the Ethics Committee of Mechnikov Institute of Microbiology and Immunology of National Academy of Medical Sciences (protocol № 4-2022 of 14.12.2022). License for the software system Yasara # 179465823.

Author Contributions

All authors made a significant contribution to the work reported, whether that is in the conception, study design, execution, acquisition of data, analysis and interpretation, or in all these areas; took part in drafting, revising or critically reviewing the article; gave final approval of the version to be published; have agreed on the journal to which the article has been submitted; and agree to be accountable for all aspects of the work.

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Disclosure

The authors declare no conflicts of interest in this work.

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