

Proceeding Paper

Application and Synthesis of a New Hybrid Heterocyclic Derivative with Antioxidant Evaluation [†]

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Abstract: New hydrazinecarboximidamide derivatives were successfully synthesized with excellent yields. These compounds have been proven to be potent inhibitors of acetylcholinesterase and tubulin, and they also exhibit significant antioxidant effects. Furthermore, an ADMET study was conducted, which suggests that these compounds not only possess the potential to cross the blood-brain barrier. These findings allow us to consider this product as a promising candidate for future medicinal development, particularly in the treatment of neurodegenerative diseases and cancer.

Keywords: heterocyclic compounds; hydrazinecarboximidamide; quinoline; antioxidant activity; theoretical simulation; in silico study; bioactives compounds

1. Introduction

Hydrazine compounds have wide applications and are very interesting, including a wide range of biological activities, antibacterial [1], antidiabetic [2], antiviral [3], antimicrobial [4], anticancer [5], Anti-Alzheimer [6] and antioxidant [7]. Such properties make these compounds particularly promising for the development of treatments targeting oxidative stress-related diseases.

The presence of numerous nitrogen atoms in these compounds grants them strong chelation properties with free radicals, which in turn significantly enhances their ability to interact effectively with target molecules. Such properties make these compounds particularly promising for the development of treatments targeting oxidative stress-related diseases.

New hydrazine derivatives were synthesized using a simple condensation reaction. These compounds have shown strong inhibitory effects on acetylcholinesterase and tubulin, as well as antioxidant properties. Furthermore, ADMET analysis confirms that these compounds adhere to Lipinski's Rule of Five and can cross the blood-brain barrier without exhibiting any ADMET-related toxicity.

2. Materials and Methods

2.1. Material Used

The ligand was prepared using the Avogadro program, while target preparation was carried out in Chimera version 1.16. Energy minimization was performed using Swiss-PdbViewer (SPDBV) version 4.1, and the visualization and analysis were conducted using BIOVIA Discovery Studio 2021, DPPH analyse using ANOVA SPSS software 2.0.

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2.2. Statistical Analysis

The reported results represent the average of three measurements, expressed as mean \pm standard deviation (SD). The IC₅₀ value, indicating the concentration required for 50% inhibition, was determined for each sample by plotting the percentage of inhibition against concentration and analyzing the resulting curve using linear regression. To identify statistically significant differences ($p < 0.05$) between the samples, a one-way analysis of variance (ANOVA) was performed using SPSS software.

2.3. Methods

2.3.1. Ligand Preparation

Preparation of derivative compounds for docking studies involved to designing small molecules, optimize their geometry, and minimize their energy. The resulting structures were then saved in the .pdb format.

2.3.2. Targets Preparation

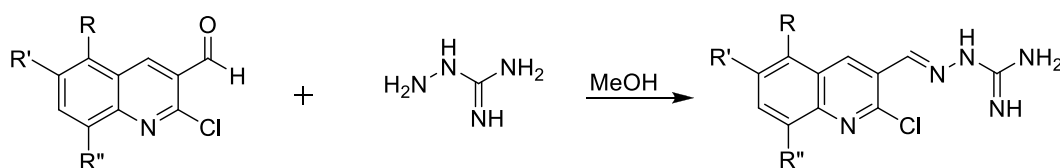
The acetylcholinesterase (AChE) protein (PDB ID: 4m0e) [8], and tubulin protein (PDB ID: 1sa0) [9], were obtained in .pdb format from the Protein Data Bank. The targets were prepared by removing water molecules, trimming unnecessary chains, and adding polar hydrogens. Energy minimization was then performed, and the results were visualized and analyzed.

2.3.3. Synthesis of Compounds

An equimolar mixture of quinoline derivatives and aminoguanidine was stirred at reflux in methanol (MeOH) without a catalyst. The precipitate was then filtered, washed with MeOH, and recrystallized from MeOH.

3. Results and Discussion

Hydrazine carboximidamides are easily synthesized through a simple condensation reaction (Scheme 1) between quinoline carbaldehyde and aminoguanidine, without the use of a catalyst. The products obtained are (E)-2-((2-chloro-5,8-dimethoxyquinolin-3-yl)methylene)hydrazinecarboximidamide and (E)-2-((2-chloro-6-methoxyquinolin-3-yl)methylene)hydrazinecarboximidamide, with yields ranging from 93% to 98%. These compounds are identified by ¹H NMR, The characteristic CHN condensation peak appears at 8.75 ppm.



A1: R=H, R'= OMe, R''=H
 A2: R=OMe, R'= H, R''=OMe

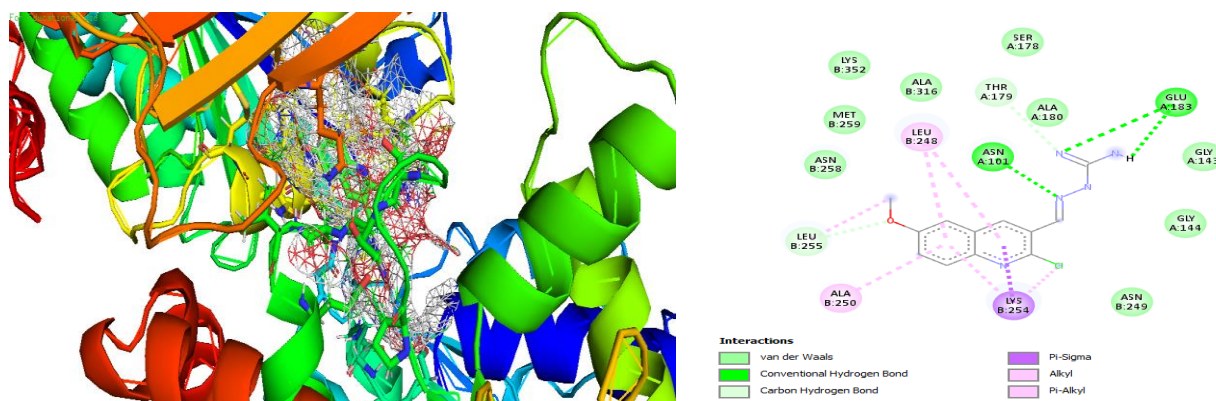
Scheme 1. General procedure for synthesis.

3.1. Docking Study

The docking study of these compounds was performed using Chimera to evaluate their inhibitory ability. In this study, the active site of the target atoms was fixed, while the chromophore remained flexible. As is well known, this program uses a genetic algorithm (GA) to dock flexible ligands into the protein binding site

The molecular docking study indicated a strong affinity and high scores within the active sites of tubulin, suggesting potential anticancer activity. However, docking with

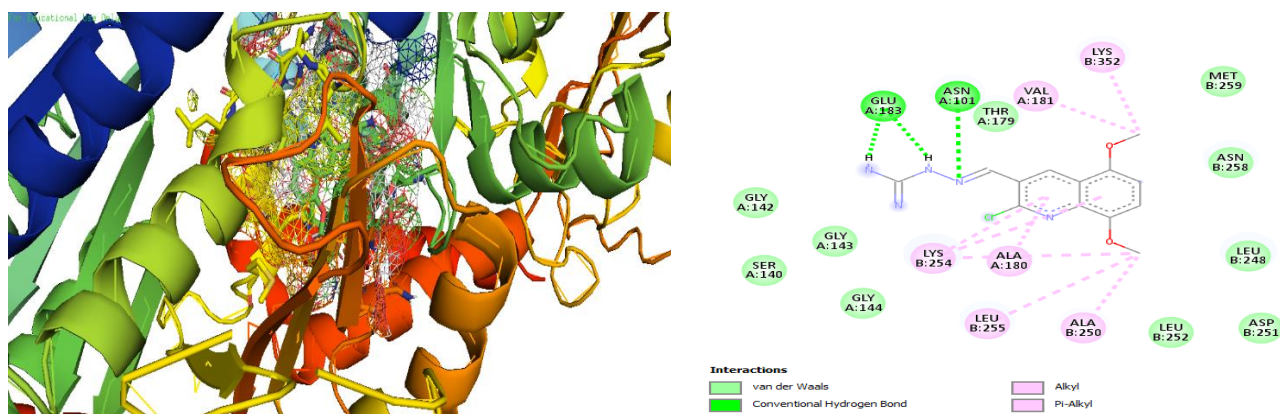
acetylcholinesterase showed a high score but with unfavorable binding, highlighting the compounds' potential for anticancer and anti-Alzheimer's activities (Figures 1–4).



A1 with Tubuline

Figure 1. Simulation of A1 in the active site of Tubuline.

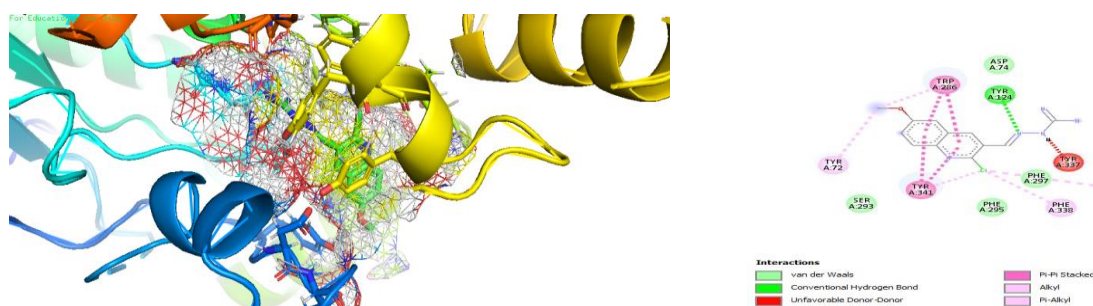
As we can see A1 is inside Tubulin (Figure 1), with good affinity due to Van der Waals interactions with ASN B:258, MET B:259, LYS B:352, ALA B:316, SER A:178, ALA A:180, GLY A:143, GLY A:144, ASN B:249, hydrogen bond GLU A:183, ASN A:101, carbon hydrogen bond THR A:179, Pi-Sigma LYS B:254, Pi-alkyl, alkyl ALA B:250, LEU B:248.



A2 with Tubuline

Figure 2. Simulation of A2 in the active site of Tubuline.

As we can see in (Figure 2), A2 is inside Tubulin with good affinity due to Van der Waals interactions with THR A:179, MET B:259, ASN B:258, LEU B:248, ASP B:251, LEU B:252, GLY A:144, SER A:140, GLY A:143, GLY A:142, hydrogen bond GLU A:183, ASN A:101, Alkyl and Pi Alkyl VAL A:181, LYS B:352, ALA B:250, LEU B:255, ALA A:180, LYS B:254.



A1 with AchE

Figure 3. Simulation of A1 in the active site of AChE.

A1 is inside AchE (Figure 3), with good affinity due to Van der Waals interactions with ASP A:74, HIS A:447, GLY A:121, SER A:203, PHE A:297, PHE A:295, SER A:293, hydrogen bond TYR A:124, Pi-Pi stacked tyr A:341, trp A:286, VAL A:294, PHE A:338, TYR A:72.



A2 with AchE

Figure 4. Simulation of A2 in the active site of AChE.

A2 is inside AchE (Figure 4), with good affinity due to Van der Waals interactions with HIS A:447, GLY A:121, SER A:203, TYR A:124, PHE A:295, ARG A:296, VAL A:294, LEU A:76, carbon hydrogen asp A:74, Pi-Pi stacked TYR A:341, TRP A:286, Alkyl TYR A:72, LEU A:289, PHE A:297, PHE A:338, unfavorable donor-donor TYR A:337.

3.2. Pharmacokinetic Study

According to the ADMET study by SwissADME (<http://www.swissadme.ch/index.php>) accessed on 6 august 2024, both A1 and A2 adhere to Lipinski's rules and Veber's rules. By ADMETSAR, both compounds are able to cross the blood-brain barrier, which suggests they may reach the brain and potentially be effective as anti-Alzheimer's agents. Additionally, we used admetSAR (<http://lmm.d.ecust.edu.cn/admetSar1>) accessed on 6 august 2024 to confirm that they are non-AMES toxic and non-carcinogenic (Table 1).

Table 1. Table of ADMET study results.

Entries	Lipinski'srules	Veber'srules	BBB	AMES Toxic	Carcinogenic	P-Glycoprotein Inhibitor
A1	+	+	+	-	-	-
A2	+	+	+	-	-	-

Significance: Yes (+); No (-).

3.3. Antioxidant Evaluation

Estimation of DPPH Free Radical Scavenging Activity

The antioxidant activity of two synthesized products, was evaluated using a modified DPPH scavenging assay [10]. Different concentrations of A1, A2, and a standard antioxidant (BHA) were prepared. Following a 30-min incubation in the dark, the absorbance was measured at 517 nm. This measurement was used to determine the ability of each compound to neutralize DPPH radicals (Table 2). The antioxidant capacity of each compound was then expressed as an IC₅₀ value (μg/mL), which represents the concen-

tration needed to reduce DPPH activity by 50% (Figure 5). Additionally, the percentage of DPPH radical inhibition for each sample was calculated using the following formula:

$$\% \text{ Inhibition} = [(A \text{ control} - A \text{ sample})/A \text{ control}] \times 100$$

where:

A control: Absorbance of the DPPH solution in methanol only.

A sample: Absorbance of the tested product or standard.

Table 2. Table of Antioxidant Evaluation.

Concentration (µg/mL)	1.5625	3.125	6.25	12.5	25	50	100	IC ₅₀ (µg/mL)
BHA	21.47 ± 0.27	34.28 ± 0.27	47.49 ± 0.27	71.09 ± 0.20	87.08 ± 0.18	89.56 ± 0.18	91.21 ± 0.10	7.18 ± 0.17
Concentration (µg/mL)	31.25	62.5	125	250	500	1000	2000	/
A1	17.94 ± 0.27	36.87 ± 0.57	43.95 ± 0.45	60.65 ± 0.27	80.29 ± 0.27	91.98 ± 0.57	92.98 ± 0.10	168.06 ± 2.92
Concentration (µg/mL)	125	250	500	1000	2000	4000	8000	/
A2	4.01 ± 0.91	7.49 ± 0.84	12.15 ± 0.67	16.87 ± 0.45	24.78 ± 0.47	63.54 ± 1.42	81.65 ± 0.62	3902.15 ± 40.35

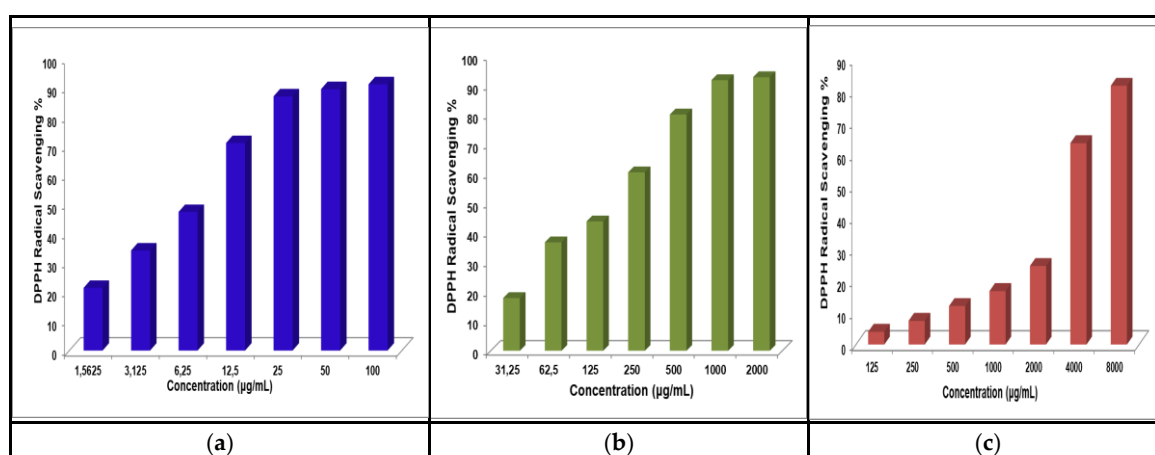


Figure 5. (a) DPPH radical scavenging activities of various concentrations of BHA standard; (b) DPPH radical scavenging activities of various concentrations of A1; (c) DPPH radical scavenging activities of various concentrations of A2.

4. Conclusions

New Hydrazine carboximidamides were successfully synthesized with excellent yields using a simple condensation reaction. These compounds demonstrated effective inhibition of tubulin and AChE in molecular docking studies. Additionally, the ADMET analysis indicates that these compounds can cross the blood-brain barrier (BBB), suggesting their potential as anti-Alzheimer's agents. Furthermore, they have proven to be antioxidant compounds, especially A1, with an IC₅₀ of 168.06 ± 2.92 µg/mL.

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