

Research article

## Synthesis and molecular docking analysis of Oxazetidine derivatives for neurological disorders

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### ABSTRACT

A series of Oxazetidine (NL1-NL12) from reacting tryptophan and aromatic aldehydes were synthesized in good yields by involving 2-[[4-chlorophenyl] methylidene] amino}-3-(1H-indol-3-yl) propionic acid and chloro acetyl chloride as reactive intermediates. All the synthesized derivatives were screened via spectral techniques. Synthesized molecules were virtually screened against Human A2A Adenosine receptor interactions analysis using molecular docking to elucidate CNS potential. Synthesized derivatives showed excellent binding towards the Human A2A Adenosine receptor.

**Keywords:** Oxazetidine, Human A2A Adenosine receptor, Docking, CNS

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### INTRODUCTION

Azetidines, one of the important heterocyclic chemical classes which are extensively researched in the scientific world. Chemically the azetidines is a four-member ring structure that contains nitrogen as heteroatom [1-2]. Importance of the azetidines can be understood by the fact it is a constituent of a number of the key chemical structures, raw materials, and important catalysts. Biologically the azetidines class of molecules is active showing a variety of therapeutic potentials like antimicrobial, anticancer, anti-inflammatory, antidepressant, antitubercular, antimalarial, anticancer, antiviral, antioxidant, and cardiovascular activities [3-11]. In recent literature the physical properties of the azetidines are reported this heterocycle is known as a conformationally puckered rigid structure, having bond angles of 10-20 degrees which depends on the type of substitution which is present on it [1-2]. Due to its attractive nature and diverse physicochemical property space associated with it, the azetidines nucleus has been extensively researched and various chemical methodologies were reported for synthesis. The chemical nature and ring strain associated with it the azetidines have become one of the most difficult chemical scaffolds to prepare, due to this reason very few methods have been successfully applied for the development of a diverse set of azetidines derivatives. Methods like Ring Closure

by C-N Bond Formation, Ring Closure by C-C Bond Formation, Reduction of  $\beta$ -Lactams, and Cyclo addition Reactions have been the most utilized chemical reactions for the synthesis of the azetidines [12]. Azetidines nucleus is not only biologically useful but its chemical applications can be widely observed in the chemical literature. In reactions like ring-opening to acyclic amines, ring expansion to pyrrolidines and another 5-Membered heterocycle, ring expansion to six-member Heterocycles, ring expansion to medium-sized heterocycle azetidines are commonly utilized. Adenosine is one of the important biochemical components which is present in the human body it can be called nucleotides which is one of the important building blocks of the genetic material in the human being. The activity of the adenosine is under the control of the producing/degrading enzymes and receptors like A1, A2A, A2B, and A3 which are specialized receptors that can be classified under G protein-coupled receptors [13-14]. These adenosine receptors are targeted for several ailments, recently the role of the A2A receptor has been proven in neurodegenerative diseases. This research paper reports development of novel Oxazetidine derivatives and their in-silico analysis for potential Adenosine A2A receptor inhibition. 12 different Oxazetidine derivatives have been synthesized and screened in silico

for potential Adenosine A2A receptor inhibition via molecular docking. All the synthesized derivatives showed excellent binding potentials towards adenosine A2A receptor which indicated these scaffolds can be utilized for the development of potent molecules targeting neurodegenerative diseases.

### EXPERIMENTAL

Synthesis of the targeted Oxazetidine derivatives was achieved in two steps as shown in figure no1 and table no1

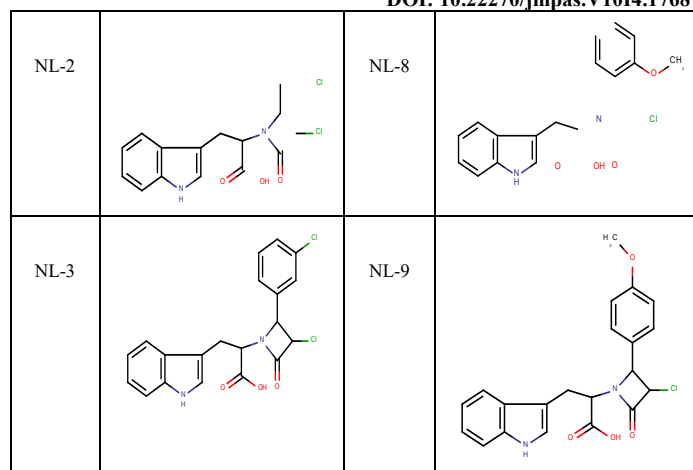
#### Step 1: Synthesis of 2-[(4-chlorophenyl) methylene] amino}-3-(1H-indol-3-yl) propionic acid

0.01M tryptophan mixed with 0.02M Benz aldehyde in 45 ml methanol and 30 ml water (60:40 ratio). This solution refluxed for 12 hours. After the reflux solution was cooled and the solvent was evaporated. The residue (reaction product) was collected and is recrystallized from methanol.

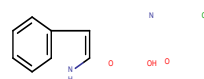
#### Step 2: Synthesis of Oxazetidine derivatives (NL1-NL12)

0.02M step-I product dissolved in 30 ml dioxane then stirred for 10 minutes in magnetic stirrer. After 10 minutes of stirring added 0.025M chloroacetyl chloride by using a dropping funnel and again stirred for 10 minutes. After 10 minutes of stirring added 0.025M triethylamine by using a dropping funnel and the solution stirred for 04 hours. After completion of stirring, the solution

Figure 1: Scheme of Synthesis of the designed Molecules

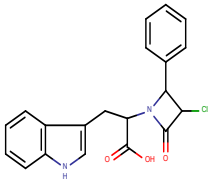
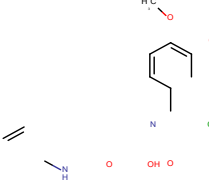


NL-4



Refluxed for 18 hours. After reflux, the final solution was poured in ice-cold water, and the formation of aqueous solution extract 02 times with 25ml ethyl acetate and then take ethyl acetate layer and evaporated and product was recrystallized from methanol.

Table 1: Synthesized Derivatives

Code	Structure	Code	Structure
NL-1		NL-7	

chloroacetyl chloride were carried out to get Oxazetidine derivatives.

The formation of the derivatives was ascertained by the thin layer chromatography, melting point, and various spectral techniques.

NL1:

IR (ATR, cm-1)- 3361.68 (NH stretching) , 2923.70 (COOH Stretching) , 1622.82 (C=O Stretching) , 1230.98 (C-N stretching) ,

NL2:

IR (ATR, cm-1)- 3361.68 (NH stretching) , 2917.23 (COOH Stretching) , 1668.15 (C=O Stretching) , 1230.98 (C-N stretching) , 744.02 (Ar-Cl)

NL3:

IR (ATR, cm-1)- 3365.68 (NH stretching) , 2912.23 (COOH Streaching) , 1657.30 (C=O Stretching) , 1232.22 (C-N stretching) , 741.30 (Ar-Cl)

NL4:

IR (ATR, cm-1)- 3354.09(N-H stretching) , 2923.70 (COOH Stretching) , 1622.82 (C=O Stretching) , 1231.81 (C-N stretching) , 744.93 (Ar-Cl).

<sup>1</sup>H NMR :([D,<sub>6</sub>]DMSO): δ 3.27(s,3H-CH<sub>3</sub>), 3.32 (s, 2H,-CH<sub>2</sub>), 3.32 (s, 2H,-CH<sub>2</sub>),3.39 (s, 2H,-CH),3.42 (s, 2H,-CH),3.42 (s, 2H,-CH<sub>2</sub>),7.04-7.09 (d, 2H, Ar-H), 7.01-7.05(d, 2H, Ar-H),10.89(s, 1H,NH),12.93(s, 1H,OH).

<sup>13</sup>C NMR ([D,<sub>6</sub>] DMSO): δ = 54.72 (-CH<sub>2</sub>), 27.14(-CH<sub>3</sub>)118.27-136.33 (Ar-C), 109.63(=CH) 160.20(N=C) 169.90 (C=O).

EIMS (M/z) : Molecular weight Correspond to 311.1molecular ion peak

NL5:

IR (ATR, cm-1)- 3368. 83 (N-H stretching), 2984. 89 (COOH Stretching), 1648.91 (C=O Stretching), 1231.10 (C-N stretching), 746. 82 (Ar-Cl)).

NL6:

IR (ATR, cm-1)- 3344.08 (N-H stretching), 2917.29 (COOH Stretching) , 1626.45 (C=O Stretching) , 1230.98 (C-N stretching) , 744.02 (Ar-OH)).

NL7:

IR (ATR, cm-1)- 3351.06 (N-H stretching), 2984.75 (COOH Stretching) , 2688.60 (Ar-OCH<sub>3</sub>), 1626.45 (C=O Stretching) ,1267.49 (C-N stretching) , 781.17 (Ar-Cl)).

NL8:

IR (ATR, cm-1)- 3344.08 (N-H stretching), 2984.65 (COOH Stretching) , 2688.60 (Ar-OCH<sub>3</sub>), 1626.45 (C=O Stretching) , 1242.69 (C-N stretching) , 743.95 (Ar-Cl)

NL9:

IR (ATR, cm-1)- 3384.40 (N-H stretching), 2922.78 (COOH Stretching) , 2682.40 (Ar-OCH<sub>3</sub>), 1652.00 (C=O Stretching) , 1341.13 (C-N stretching) , 742.53 (Ar-Cl).

<sup>1</sup>H NMR :([D,<sub>6</sub>]DMSO): δ 3.56(s,H-CH<sub>3</sub>), 3.08(s,H-CH<sub>2</sub>), 3.10 (s, H,-CH<sub>2</sub>), 4.08 (s, H,-CH), 4.48 (s, H,-CH), 4.50 (s, H,-CH), 6.90-7.00 (m, Ar-H), 7.13-8.48(m, Ar-H),10.88(s, 1H,NH),12.83(s, 1H,OH),

EIMS (M/z) : Molecular weight Correspond to 380.2 molecular ion peak

NL10:

IR (ATR, cm-1)- 3260.08 (N-H stretching), 2928.58 (COOH Stretching) , 1627.45 (C=O Stretching) , 1347.13 (C-N stretching) , 731. 85 (Ar-Cl).

<sup>1</sup>H NMR :([D,<sub>6</sub>]DMSO): δ 3.18(s,H-CH<sub>2</sub>), 3.20 (s, H,-CH<sub>2</sub>), 4.08 (s, H,-CH), 4.48 (s, H,-CH), 4.50 (s, H,-CH), 6.95-7.00 (m, Ar-H), 7.13-8.48(m, Ar-H),10.88(s, 1H,NH),12.83(s, 1H,OH),

EIMS (M/z) : Molecular weight Correspond to 397.3 molecular ion peak

NL11:

IR (ATR, cm-1) - 3368.83 (N-H stretching), 2917.73 (COOH Stretching), 1656.11 (C=O Stretching), 1231. 81 (C-N stretching), 740. 89 (Ar-Cl).

NL12:

IR (ATR, cm-1)- 3344.08 (N-H stretching), 2984.65 (COOH Stretching), 1626.45 (C=O Stretching) , 1233.55 (C-N stretching), 747.40 (Ar-Cl), 555.71 (Ar-Br).

### Molecular Docking

Molecular docking analysis was utilized to predict the possible binding mode of the developed molecules against the human adenosine A<sub>2A</sub> receptor. Docking analysis revealed that all the developed molecules have a good binding ability with the human adenosine A<sub>2A</sub> receptor. Derivative NL-1 Showed hydrogen bond interaction with ILE66and aromatic interactions with ILE66, hydrophobic interactions with LEU167 as shown in figure no2. Derivative NL-2 Showed hydrogen bond interaction with SER67and aromatic interactions with HIS264, TYR271, hydrophobic interaction with LEU167 as shown in figure no 3

Figure 2: Docking Interactions of derivative NL1

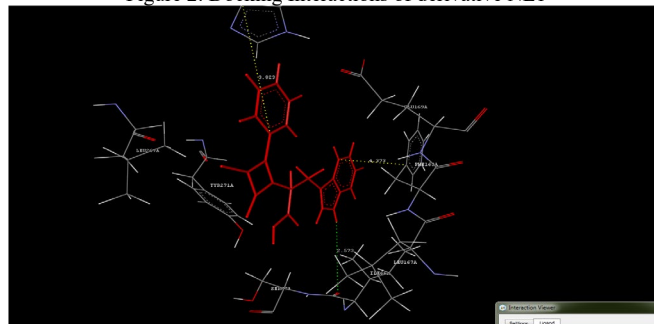
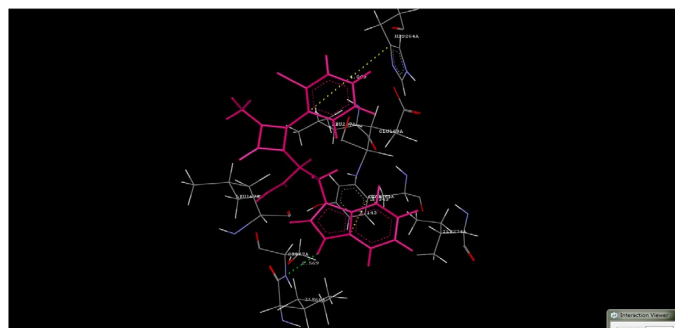
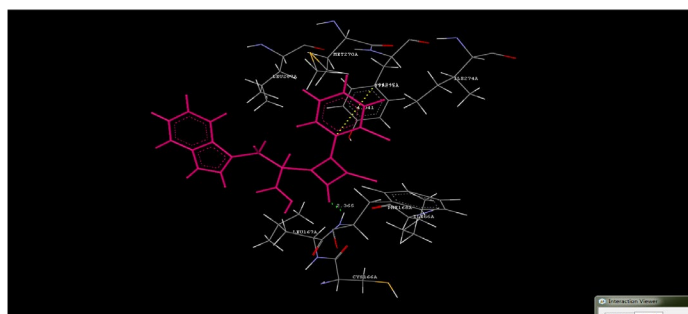


Figure 3: Docking Interactions of derivative NL2



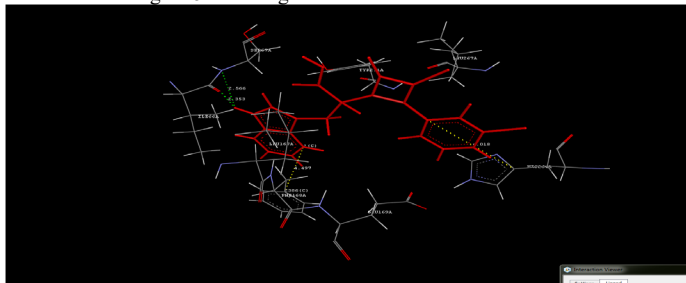
Derivative NL-3 Showed hydrogen bond interaction with PHE168and aromatic interactions with TYR271, hydrophobic interaction with LEU167, LEU267 as shown in figure no 4.

Figure 4: Docking Interactions of derivative NL3



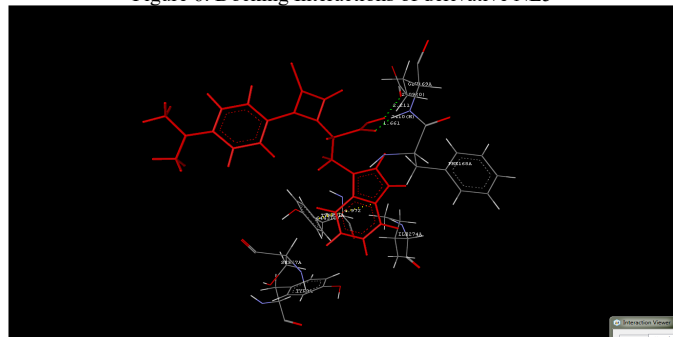
Derivative NL-4 Showed hydrogen bond interaction with

Figure 5: Docking Interactions of derivative NL4



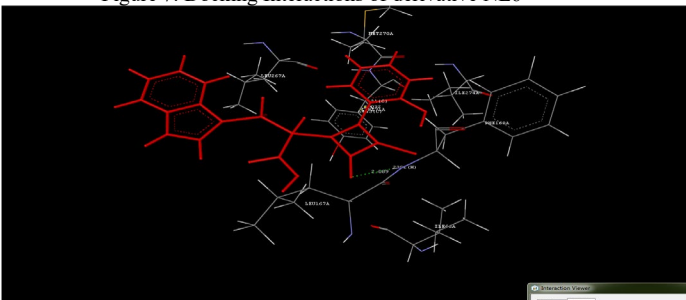
Derivative NL-5 Showed hydrogen bond interaction with GLU169 and aromatic interactions with TYR271 as shown in figure no 6.

Figure 6: Docking Interactions of derivative NL5



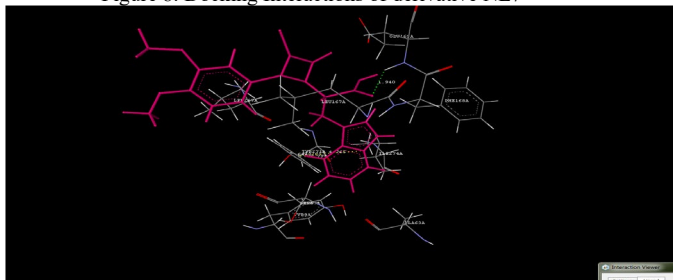
Derivative NL-6 Showed hydrogen bond interaction with PHE168 and aromatic interactions with TYR271 has shown in figure no 7.

Figure 7: Docking Interactions of derivative NL6



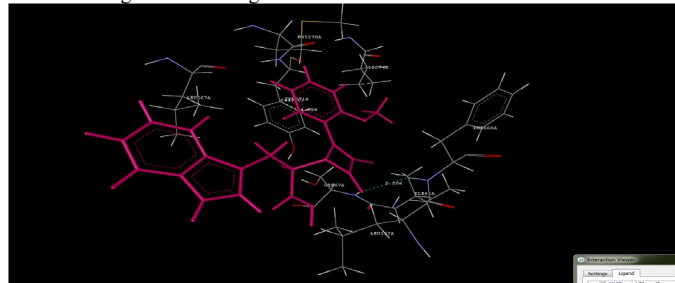
Derivative NL-7 Showed hydrogen bond interaction with GLU169 and aromatic interactions with TYR271 has shown in figure no 8.

Figure 8: Docking Interactions of derivative NL7



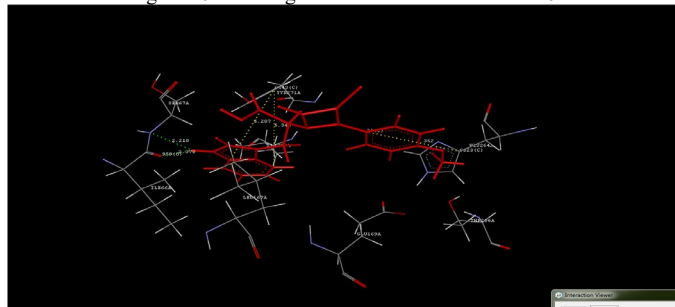
Derivative NL-8 Showed hydrogen bond interaction with PHE168 and aromatic interactions with TYR271 has shown in figure no 9.

Figure 9: Docking Interactions of derivative NL8



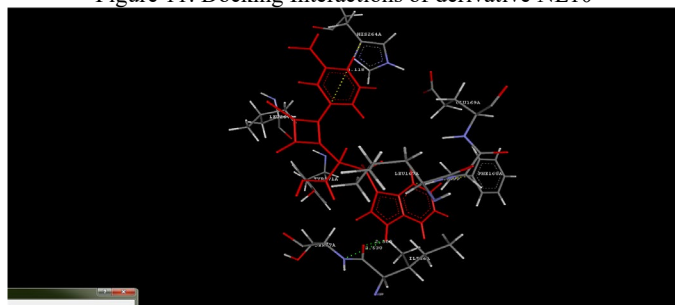
Derivative NL-9 Showed hydrogen bond interaction with SER67, ILE66 and aromatic interactions with HIS264, TYR271 has shown in figure no 10.

Figure 10: Docking Interactions of derivative NL9



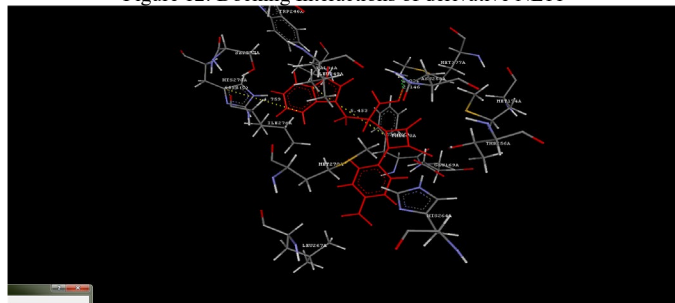
Derivative NL-10 Showed hydrogen bond interaction with SER67, ILE66 and aromatic interactions with PHE168, HIS264 has shown in figure no 11

Figure 11: Docking Interactions of derivative NL10



Derivative NL-11 Showed hydrogen bond interaction with ASN253 and aromatic interactions with PHE168, HIS278 has shown in figure no 12

Figure 12: Docking Interactions of derivative NL11



Derivative NL-12 Showed hydrogen bond interactions with has shown in figure no 13.

Figure 13: Docking Interactions of derivative NL2



## CONCLUSION

New Oxazetidine derivatives incorporating various heterocyclic Benzaldehyde substituents are prepared by using 2-[[4-chlorophenyl] methylidene] amino}-3-(1H-indol-3-yl) propionic acid and chloro acetyl chloride as reactive intermediates. All these derivatives are characterized by spectral data. All the synthesized products are screened for their in silico Human A2A binding analysis via docking analysis. The majority of the tested compounds exhibited significant Human A2A binding ability. Hence this study has widened the scope of development potent molecules targeting neurological diseases.

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