

Molecular docking, elucidating the regiospecificity and the mechanism of [3+2] cycloaddition reaction between azidobenzene and propionaldehyde

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ABSTRACT

Molecular electron density theory has been performed with the B3LYP/6-31(d,p) method to study the [3+2] cycloaddition processes between azidobenzene and propionaldehyde, the reactivity indices, activation and reaction energies are computed. The reaction and activation energies indicate that this [3+2] cycloaddition reaction is regiospecific, in good agreement with the experimental results. ELF examination revealed that the mechanism of these cycloaddition reactions takes place in two steps. In addition, a docking approach was performed on the products investigated, and the interaction with the protein protease COVID-19 (PDB ID: 6LU7), the results confirm that the presence of triazole and isoxazole rings increases the affinity of these products.

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1. Introduction

Triazoles are five-membered heterocyclic compounds containing three nitrogen atoms and two carbon atoms¹ (Fig. 1). Triazoles are often used as ligands in coordination chemistry and catalysis, as well as in organic synthesis for the formation of carbon-nitrogen bonds and are often synthesized using the nitrogen-alkyne cycloaddition reaction² and are also present in some drugs and in compounds with interesting biological properties, such as antimicrobial, antiviral and anticancer properties³. These properties have led to studies on the potential of triazole as a drug candidate in the treatment of various diseases. Another family of heterocyclic organic compounds is the isoxazoles, which have one oxygen atom and one nitrogen atom in the ring (Fig. 1). 2-isoxazoline has applications in pharmaceutical and agrochemical chemistry⁴. They can be used as precursors in the synthesis of other heterocyclic compounds and are used as herbicides and fungicides in agriculture, they also have applications in the synthesis of drugs and pharmaceuticals, and have been studied for their analgesic, anti-inflammatory and antinociceptive properties⁵, isoxazoles can be prepared by cycloaddition reactions between nitriloxides and alkenes⁶ (Fig. 1).

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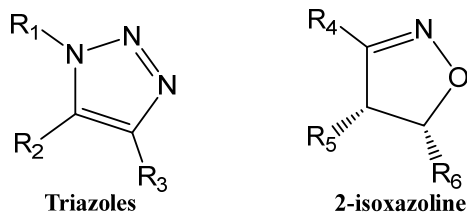
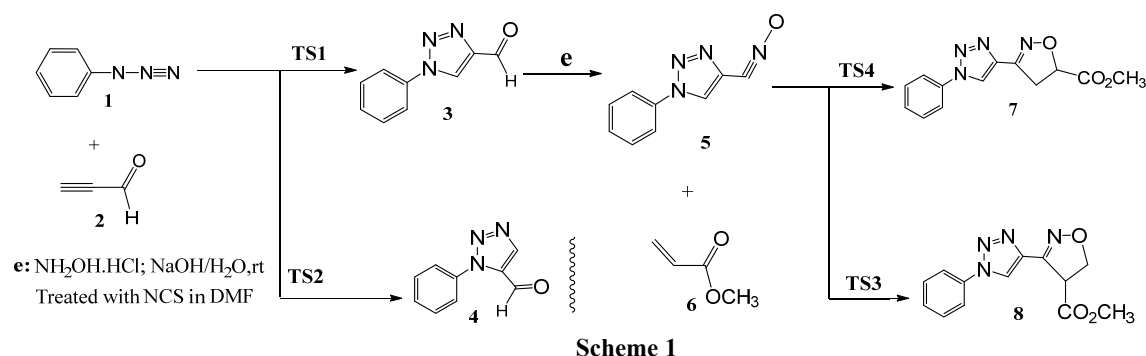


Fig. 1. Triazoles and isoxazoles structures

The computerized method known as DFT (Density Functional Theory) in physics and chemistry is used to characterize the electrical properties of molecules, materials, and surfaces⁷. The strategy was created using the quantum theory of matter, which characterizes subatomic particles in terms of waves and wave functions. Within the framework of the DFT method another theory has recently been proposed called MEDT, which is very good at predicting the electrical and physical properties of the chemical structures. It is commonly used for modeling chemical reactions in addition to predicting chemical reactivity⁸. This concept has been applied in several surveys to explain how reactive cycloaddition mechanisms work⁹⁻²⁰.

In order to understand the processes and the selectivity of the cycloadditions reactions mentioned in **Scheme 1**, the MEDT has been used technique to explore the selectivity and mechanism of these [3+2] cycloadditions reactions²¹.



2. Computation Methodology

B3LYP is a functional approach in DFT theory which has been applied to describing the electrical and energetic properties of molecular structures²². Basis 6-311(d,p) has been applied to study electron distribution in molecules²³. All calculation processes are performed in Gaussian 09²⁴. The transition states (TS) were identified and verified through the existence of a singular imaginary frequency. The IRC also helps to evaluate the reaction trajectory from its initial to final state, and to understand the intermediate steps and the activation barriers attached to them. Tomas' PCM has been used to take the solvent's impact on the optimized structures into account. (Polarizable Continuum Model)²⁵⁻³⁰. The electronic chemical potential is a thermodynamic measure that indicates the tendency of a reactant to gain or lose electrons from its electronic structure, and it is described as follows: $\mu = \frac{E_L + E_H}{2}$. Chemical hardness is a concept from chemistry describing a molecule's resistance to disorder in its electronic structure, which is defined by the relationship: $\eta = E_L - E_H$, based on the energies of the HOMO and LUMO boundary molecular orbitals, expressed as: E_H and E_L .

The measure designated global electrophile index (ω) is a quantity used in chemistry to appreciate the capacity of a molecule to intervene as an electrophile, in other terms to capture electrons from other molecules to establish a chemical bond, and is explained by the formula below: $\omega = \frac{\mu^2}{2\eta}$ ³¹. Furthermore, in opposite to the electrophile index, the nucleophile index (N) is a useful measure of a molecule's capacity to behave as a nucleophile, i.e. to donate electrons to another molecule to create a chemical bond, which is given as follows: $E = E_{\text{HOMO}}(\text{Nu}) - E_{\text{HOMO}}(\text{TCE})$ ³².

Parr functions are used to calculate local indices, which are fundamental spin functions in quantum chemistry to describe the distribution of electron spins in molecules^{33,34}. The electron localization function is based on the Topmod program (ELF)^{35,35}.

3. Results and discussion

3.1. Analysis of the CDFT indices for reagents

Numerous investigations on chemical reactions²⁵⁻³⁰ have shown that evaluating the reactivity indices provided inside CDFT is an efficient method for assessing the reactivity of organic molecules. In order to forecast the reactivity of phenylazide (**1**), propionaldehyde (**2**), nitrile-oxide (**5**) and methyl-acrylate (**6**) in the cycloaddition processes, the global indices electrophile, nucleophile, chemical hardness, and electronic chemical potential are explored in **Table 1**.

Table 1. Values of phenylazide (**1**), propionaldehyde (**2**), nitrile-oxide (**5**) and methyl-acrylate (**6**), as determined by B3LYP/6-31G(d), in terms of nucleophile N, electrophile, electrochemical potential, and chemical hardness; all values are in electron volts (eV).

System	η	μ	ω	N
1	5.16	-3.62	1.26	3.32
2	5.53	-4.80	2.08	1.96
5	4.91	-4.00	1.62	3.07
6	6.16	-4.31	1.50	2.13

While the disparity in chemical electron potentials suggests³⁷ that this cycloaddition reaction is polar, the chemical electron potential of **1** ($\mu = -3.62$ eV) is lower than that of **2** ($\mu = -4.80$ eV), suggesting to an electro-flow from **1** to **2**, on the other hand **5**'s chemical electron potential ($\mu = -4.00$ eV) is lower than that of **6**'s ($\mu = -4.31$ eV), reveals that there is an electro-flow from **5** to **6**, although the difference in chemical electron potential indicates that this cycloaddition reaction between **5** and **6** is non-polar. The electrophile index ω ^{38,39} and nucleophile index N³⁵ of **1** (1.26 and 3.32 eV) and for **5** are (1.62 and 3.07 eV) respectively, and therefore its classification in the families of strong electrophiles and strong nucleophiles. The electrophile index ω of **2** and **6** are (2.08, 1.50 eV) respectively, which classifies it as a strong electrophile, and the nucleophile index N are (1.96, 2.13 eV), respectively, which classifies it as a moderate nucleophile according to the standard electrophile³⁹ and nucleophile scales⁴⁰, Inferring that whereas **2** and **6** will act as electrophiles in this reaction, **1** and **5** will participate as nucleophiles.

3.2. Local indices

Since their introduction by Luis Domingo⁴¹, Parr functions are used extensively in theoretical chemistry to predict the chemical reactivity of organic compounds. Several studies show that first bonding is established on the center with the highest Parr electrophylic value in dienophiles. Figure 2 displays the electrophilic Parr functions of the two dienophiles, **2** and **6**.

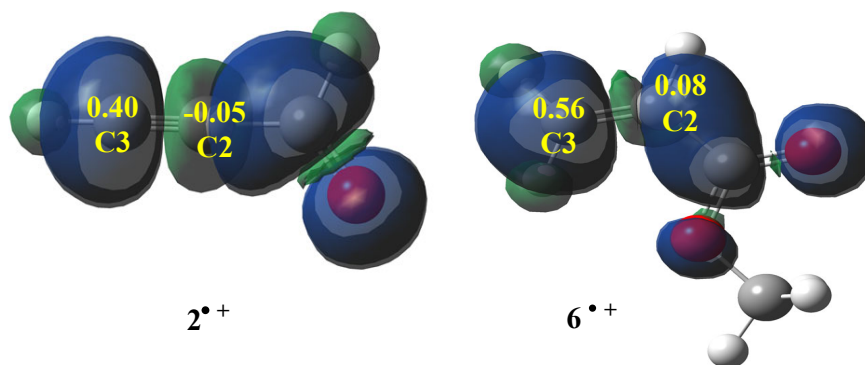


Fig. 2. The electrophilic Parr functions with 3-dimensional atomic spin density (ASP) mappings of the reagents **2** and **6**.

Fig. 2 reveals that the Mulliken atomic spin densities of propionaldehyde (**2**) $P_3^+ = 0.56$ and methyl-acrylate (**6**) $P_3^+ = 0.40$, are highly localised on the C3 centre of both molecules, while there is almost no surface area around the C2 centre. This result indicates that the C3 center is the most reactive, so the first bond will form on the C3 center of propionaldehyde (**2**) and methyl-acrylate (**6**).

4. ELF studies of reagents

The Electron Localization Function (ELF) is a technique used in quantum chemistry and solid-state physics to analyze the electronic structure of molecules. The ELF is a function that provides information about the degree of electron localization within a given region of space. The ELF can be used to identify the nature of chemical bonds within a molecule. For example, a high ELF value in a certain region indicates the presence of a covalent bond, while a low ELF value indicates

the absence of a bond or the presence of a weakly interacting species. The ELF basins for the reactants and the attractor are assigned in **Fig. 3** at their important positions.

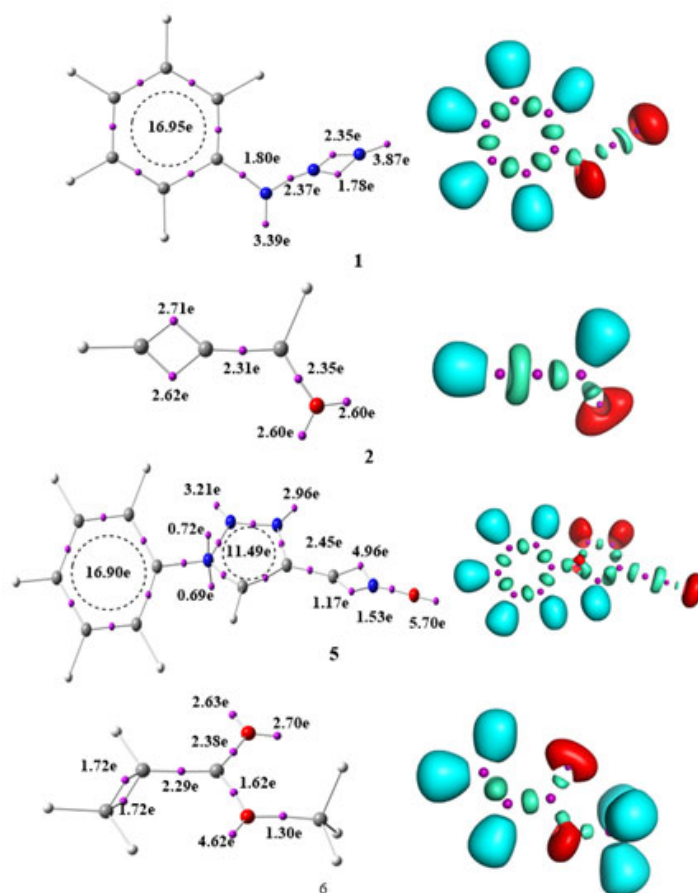


Fig. 3. B3LYP/6-31G(d,p), ELF positions basin attractors with ELF localization domains with the greatest relevance for valence basin populations are described at an ELF isosurface value of 0.80, of phenylazide (**1**), propionaldehyde (**2**), nitrile oxide (**5**) and methyl-acrylate (**6**), the average number of electrons.

The two free pairs of non-bonding electrons on the N1 and N3 nitrogen atoms are shown to be associated with a single monosynaptic basin $V(N1)$ and $V(N3)$ that has an electron population of 3.39e and 3.87e, respectively, in the phenylazide (**1**) ELF building design. The existence of two disynaptic clusters $V(N2,N3)$ and $V'(N2,N3)$ with a cumulative electron population of 4.13e indicates the presence of an $N2=N3$ double bond, while the $N1-N2$ bonding region exhibits a significant double bond nature due to the presence of a disynaptic cluster $V(N1,N2)$ with an electron population of 2.13e. Because of the 37e electron population, this structure suggests that BAZ 1 participates in the cycloaddition reaction between phenylazide (**1**) and propionaldehyde (**2**) in the form of a pseudodiradical.

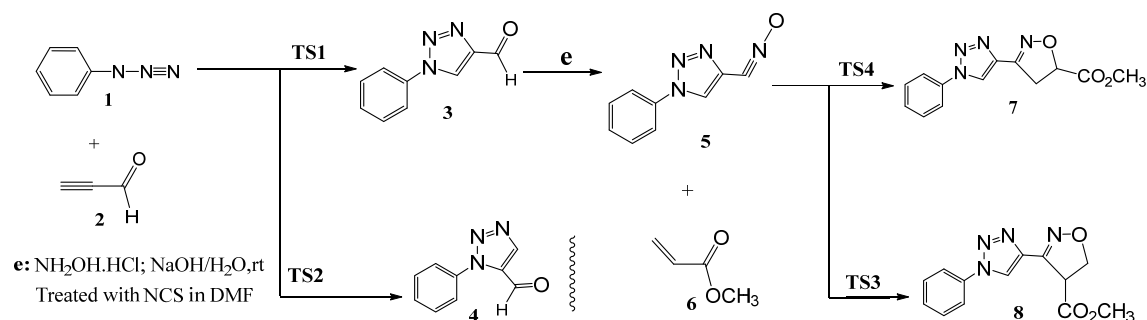
It can be observed in propionaldehyde (**2**), existence of two disynaptic pools $V(C2,C3)$ and $V'(C2,C3)$ which have a total electron population of 5.33e, one disynaptic pool $V(C1,C2)$ integrating 2.31e and existence of two monosynaptics carried by the oxygen atom of value 5.20e.

In molecule **6** we observe essentially two bisynaptic basins between carbons C2 and C3 integrating in total $VT(C2,C3)=3.42e$, in reagent **5**, we observe the presence of a monosynaptic basin around the oxygen atom of value 5.70e and two bisynaptic basins between carbon C1 and the nitrogen atom of total value 6.13e, which shows the existence of a triple bond between the carbon C1 and the nitrogen atom, another bisynaptic basin located between the nitrogen atom and the oxygen atom with a value of 1.53e, this electronic distribution clearly shows that nitrile oxide will participate in the cycloaddition reaction between nitrile-oxide (**5**) and methyl-acrylate (**6**) as a zwitterionic type.

5. Energetical considerations

Phenylazide (**1**), propionaldehyde (**2**), nitrile oxide (**5**) and methyl acrylate (**6**) are non-symmetrical reactants, so the [3+2] cycloaddition reaction between phenylazide (**1**) and propionaldehyde (**2**) will lead to two pathways and the [3+2] cycloaddition reaction between nitrile oxide (**5**) and methyl acrylate (**6**) will provide two other pathways, so in this study

there are four pathways in total (Scheme 2). The thermodynamic values are providing in table 2 and the geometries of the transitions states are displayed in Figure 4.



Scheme 2 Competitive reaction pathways involved in the 32CA reaction between 1 and 2 [5+6].

Table 2. B3LYP/6-31(d,p) relative (ΔE , ΔH and ΔG , in kcal·mol⁻¹) energies, in DMSO, for the species involved in the opening of the cycloaddition reaction between 1+2 and between 5+6.

System	ΔE	ΔH	ΔG	ΔS
1	-----	-----	-----	-----
2	-----	-----	-----	-----
1+2	-----	-----	-----	-----
TS1	15.5	15.9	27.6	-39.268
3	-73.8	-70.0	-55.3	-48.989
TS2	16.6	16.9	29.1	-40.843
4	-70.5	-66.7	-52.1	-48.939
5	-----	-----	-----	-----
6	-----	-----	-----	-----
5+6	-----	-----	-----	-----
TS3	12.8	13.2	26.5	-44.527
8	-38.5	-35.4	-21.4	-46.981
TS4	11.1	11.5	23.6	-40.566
7	-39.1	-36.3	-48.8	-46.982

Analysis of the relative energies of the stationary points involved in the 32 CA reaction between phenylazide (1) and propionaldehyde (2), leads to some interesting conclusions: (i) the most favorable TS1 is found at 15.5 Kcal·mol⁻¹ below the separate reactants and TS2 is found at 16.6 kcal·mol⁻¹ below the separate reactants; (ii) this 32CA reaction is highly exothermic at 73.8 Kcal·mol⁻¹ for product 3 and 70.5 for product 4. Therefore, the formation of product 3 occurs by kinetic and thermodynamic control. (iii) this 32CA reaction is completely ortho-regioselective, as TS2 is found to be 1.1 Kcal·mol⁻¹ higher than TS1.

Some intriguing findings are drawn from an analysis of the relative energy of the stationary points involved in the 32CA reaction between nitrile-oxide (5) and methyl-acrylate (6):

- (i) The preferable TS4 appears at 11.1 Kcal·mol⁻¹ under the separate reactant and the most favorable TS3 is discovered at 12.8 Kcal·mol⁻¹ below the separate reactants.
- (ii) (ii) this 32CA reaction is exothermic at 39.1 Kcal·mol⁻¹ for product 7 and 38.5 for product 8. Therefore, kinetic and thermodynamic control governs the creation of product 3.
- (iii) (iii) Given that TS4 has been found to be 1.1 Kcal·mol⁻¹ bigger than TS3, this 32CA reaction is entirely ortho-regioselective.

The geometries of the four TSs are illustrated in Figure 4. At the most favourable ortho TSs TS1 and TS4, the distances between the N1-C3 and C2-N3 interacting centers are 2.031 and 2.356 Å at TS1 and 2.171 and 2.439 Å at TS4, while at the meta TSs, the distances between the two interacting centres C3-N3 and C2-N1 are: 2.060 and 2.314 Å, respectively, at TS2 and 2.190 and 2.285 Å, respectively, at TS3.

These distances indicate that the most favourable TS1 and TS4 have high geometric asynchronism respectively $\Delta l=0.324$ and $\Delta l=0.268$ Å, the most unfavourable TS2 and TS3 have very low geometric asynchronism respectively $\Delta l=0.254$ and $\Delta l=0.095$ Å⁴².

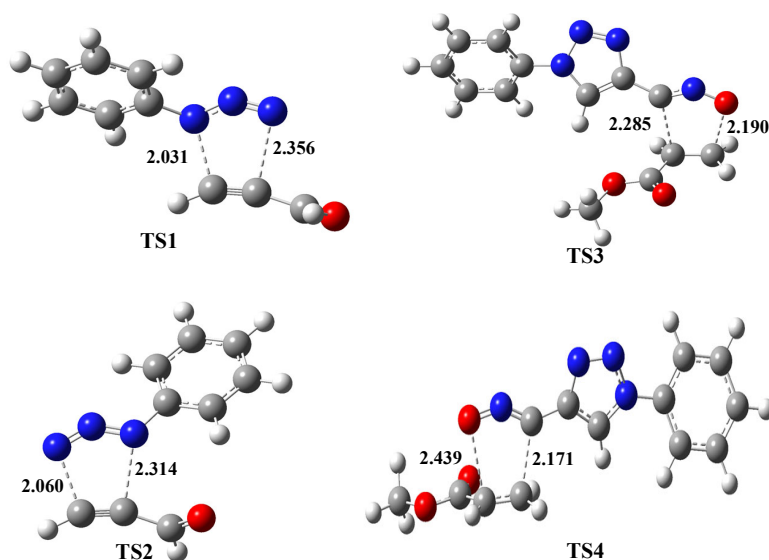


Fig. 4. B3LYP/6-31G(d,p) geometries of the optimized transition states associated with the 32CA reactions between 1 and 2 (5 and 6), distances are given in Angstrom, Å.

6. Bond evolution theory study

The mechanism of a reaction can be investigated using an electron localization function (ELF). We can track variations in electron density (electron population) along the chemical route by applying ELF to the IRC profile. This enables a thorough mechanism analysis of the reaction by allowing us to determine precisely when a given bond is being produced or broken and to investigate the interchange of the electron population in lone pair atoms and in bonds.

6.1. Mechanism of the cycloaddition reaction between phenylazide (1) and propionaldehyde (2)

Only significant locations where noticeable changes in the electron population and valence basins occur are taken into account in this context's ELF analysis of a number of chosen points on the IRC of TS1's most favorable points (**Fig. 5**), the values of the basins are compiled in **Table 3** (AZ-1 to AZ-9).

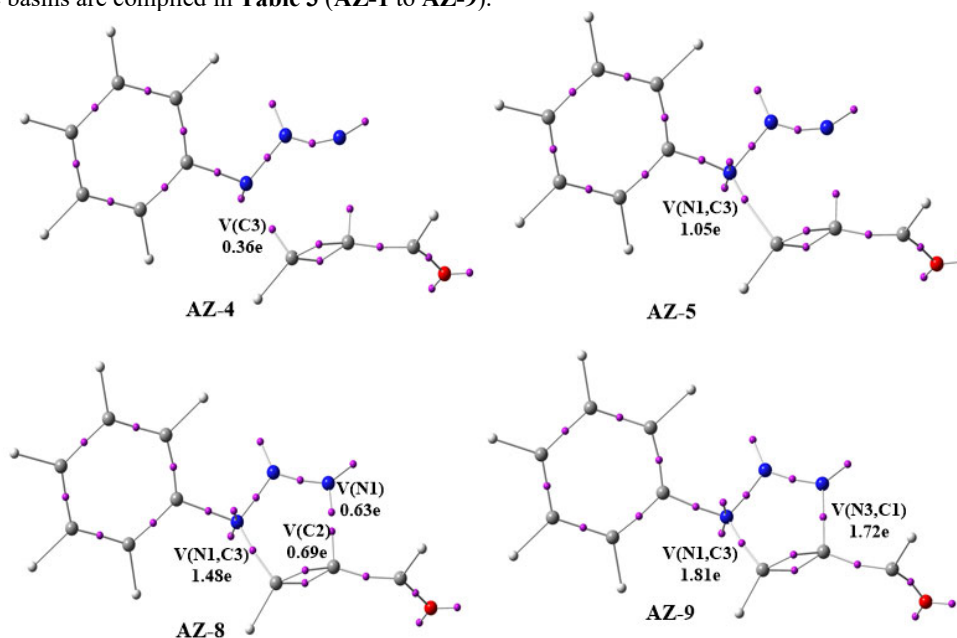


Fig. 5. Positions of the ELF valence attractors with populations for the AZ-4, AZ-5, AZ-8 and AZ-9 points involved in the IRC profile of TS-1

Table 3. ELF valence basin populations, distances of the forming bonds, of TS1 and TS4, the selected structures of the IRC involved in the formation of the new single bonds along cycloaddition between phenylazide (1) with propionaldehyde (2) and cycloaddition reaction between nitrile-oxide (5) and methyl-acrylate (6). Electron populations are given in average number of electrons, e, and distances in angstroms, Å

	d(N3-C2)	d(N1-C3)	V(N1,N2)	V(N1)	V'(N1)	V(N2)	V(N2,N3)	V(N3)	V'(N3)	V(C2,C3)	V'(C2,C3)	V(C2)	V(C3)	V(N1,C3)	V(N3,C2)	***
AZ-1-TS-1	2.568	2.032	1.74	3.32	-	2.11	2.78	3.80	-	2.43	2.42	-	-	-	-	***
AZ-2	2.221	1.964	1.67	3.34	-	2.29	2.70	3.81	-	2.43	2.34	-	-	-	-	***
AZ-3	2.221	1.964	1.67	3.34	-	2.29	2.70	3.81	-	2.43	2.34	-	-	-	-	***
AZ-4	2.125	1.875	1.64	3.36	-	2.43	2.62	3.83	-	2.24	2.22	0.52	0.23	-	-	***
AZ-5	2.030	1.784	1.62	1.96	0.69	2.55	2.56	3.85	-	2.15	2.12	0.60	-	<u>1.05</u>	-	***
AZ-6	1.981	1.739	1.61	1.91	0.69	2.60	2.52	3.87	-	2.12	2.08	0.63	-	<u>1.15</u>	-	***
AZ-7	1.881	1.6529	1.60	1.81	0.68	2.70	2.45	3.53	0.38	2.07	2.02	0.67	-	<u>1.32</u>	-	***
AZ-8	1.775	1.5759	1.60	1.71	0.63	2.80	2.37	3.34	0.61	2.05	1.96	0.69	-	<u>1.48</u>	-	***
AZ-9	1.542	1.461	1.62	1.47	0.50	3.00	2.19	3.12	-	2.05	1.80	-	-	<u>1.81</u>	<u>1.72</u>	***
	d(O-C2)	d(C-C3)	V(C4,N1)	V'(C4,N1)	V(N1)	V(C4)	V'(C4)	V(N1,O1)	V(O1)	V'(O1)	V(C2,C3)	V'(C2,C3)	V(C2)	V(C3)	V(C4,C3)	V(O1,C2)
OX-1	3.069	3.138	2.87	2.62	-	0.66	-	1.53	2.92	2.78	1.70	1.69	-	-	-	-
OX-2-TS-4	2.440	2.172	1.75	1.48	1.96	1.34	-	1.41	2.82	2.79	3.17	-	-	-	-	-
OX-3	2.402	2.113	1.69	1.44	2.07	1.10	0.27	1.39	2.83	2.77	3.15	-	-	-	-	-
OX-4	2.365	2.053	1.93	1.42	2.16	0.76	0.38	1.37	2.82	2.78	2.87	-	-	0.25	-	-
OX-5	2.327	1.993	1.91	1.38	2.25	0.81	0.37	1.35	2.81	2.78	2.75	-	-	0.33	-	-
OX-6	2.288	1.934	1.89	1.37	2.32	-	0.36	1.32	2.80	2.79	2.66	-	-	-	<u>1.25</u>	-
OX-7	1.937	1.611	3.32	-	2.69	-	-	1.16	2.84	2.78	2.13	-	0.27	-	<u>1.78</u>	-
OX-8	1.807	1.573	3.28	-	2.76	-	-	1.09	2.96	2.72	2.06	-	0.32	-	<u>1.85</u>	-
OX-9	1.740	1.561	3.26	-	2.79	-	-	1.07	2.69	2.68	2.03	-	-	-	<u>1.88</u>	<u>0.67</u>
OX-10	1.607	1.540	3.22	-	2.85	-	-	1.00	2.60	2.60	1.98	-	-	-	<u>1.92</u>	<u>0.94</u>
OX-11	1.495	1.523	3.18	-	2.90	-	-	0.95	2.56	2.52	1.95	-	-	-	<u>1.95</u>	<u>1.14</u>

In the AZ-1 structure, there are three monosynaptic basins carried by the N1, N2 and N3 nitrogen atoms, with values of 3.32e, 3.11e and 3.80e respectively. Two bisynaptic basins located between (N1,N2) and (N2,N3), which respectively carry the following values: 1.74e and 2.78e, one also observes two bisynaptic basins located between the C2 and C3 carbons with a total value $V(C2,C3)=4.85e$. The two fragments, the azide and the alkene, move closer to each other, which influences the electronic distribution around the atoms, which becomes in the AZ-3 structure: $V(N1)=3.34e$, $V(N2)=2.29e$, $V(N3)=3.81e$, $V(N1,N2)=1.67e$, $V(N2,N3)=2.70e$ and $V(C2,C3)=4.77e$. In structure AZ-4 the appearance of two new monosynaptic basins carried by carbons C2 and C3 of value $V(C2)=0.52e$ and $V(C3)=0.23e$.

In the AZ-5 structure, the monosynaptic basin $V(N1)$ decomposes into three parts $V(N1)$ with a value of 1.96e, $V'(N1)=0.96e$ and the third part will be combined with $V(C3)$ to form the first N1-C3 single bond with an electron population $V(N1,C3)=1.05e$. In the AZ-7 structure, the monosynaptic basin $V(N3)$ is decomposed into two basins $V(N3)$ and $V'(N3)$ respectively with values of 3.53e and 0.38e, these values become in the AZ-4 structure: $V(N3)=0.34e$ and $V'(N3)=0.61e$. The monosynaptic pool $V'(N3)$ will share its electron density with the monosynaptic pool $V(C2)$ to form the second N3-C2 bond.

6.2. Mechanism of the cycloaddition reaction between nitrile oxide (5) and methyl acrylate (6)

The ELF analysis of certain selected points on the IRC of most favorable TS4's (Fig. 6) takes into account only major areas where noteworthy variations in the electron population and valence basins occur; the values of the basins are collected in Table 3 (OX-1 to OX-11).

During the cycloaddition reaction between nitrile oxide (5) and methyl acrylate (6), the electronic distribution in structure OX-1 is almost identical to that of the separate reactants. In structure OX-3 we have the appearance of two monosynaptic basins carried by the C4 carbon with values of $V(C4)=1.10e$ and $V'(C4)=0.27e$, these values will increase to $V(C4)=0.81e$ and $V'(C4)=0.37e$ in structure OX-5. In structure OX-4 we observe the appearance of a new monosynaptic basin carried by carbon C3 of value $V(O3)=0.25e$. In structure OX-6 the monosynaptic basins $V(C4)$ and $V(C3)$ will share their electron density to form the C3-C4 bond with an initial density $V(C3,C4)=1.25e$.

In structure OX-7 and OX-7, we have the apparition of a new monosynaptic carried by carbon C2 of value $V(C3)=0.32e$, this basin ($V(C3)$) will form the second O1-C2 bond with a part of monosynaptic basin of the oxygen atom. As a conclusion, the bonds are formed asynchronously in the reactions studied in this work.

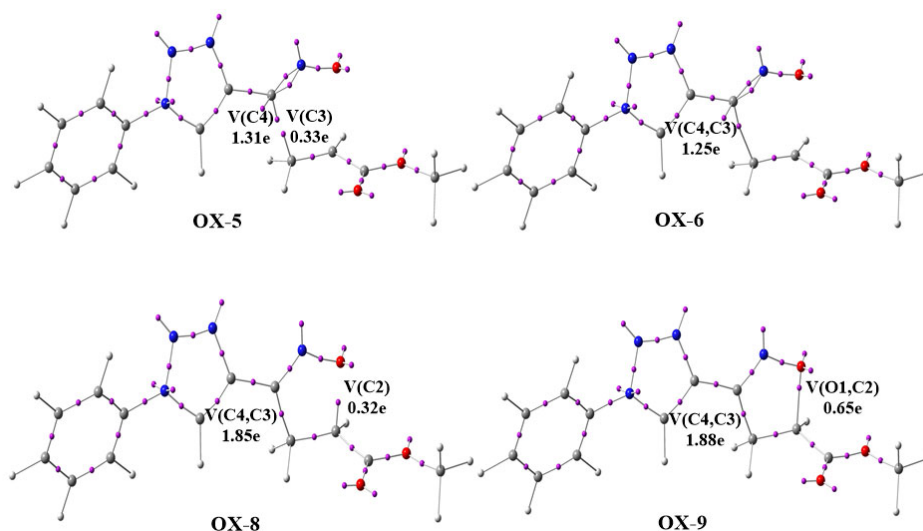


Fig. 6. Positions of the ELF valence attractors with populations for the OX-5, OX-6, OX-8 and OX-9 points involved in the IRC profile of TS-4

7. Docking analysis

Molecular docking is a modeling tool used in chemistry for researching at the relationships between ligands and proteins. In function to their chemical structure and energy potential, it allows for the prediction of the three-dimensional conformation of a molecular complex between a ligand and protein. It also enables the prediction of the affinity between a protein and a ligand, which can be helpful in the process of discovering drugs⁴³. This section compares the antiviral activity of ribavirin to that of the antiviral products **3**, **4**, **7**, and **8**, which have been studied in relation to the crystal structures of the SARS-CoV-2 protease or the main COVID-19 protease (Mpro) in access numbers 6LU7. Biovia's Discover Studio 3.5 and PyMol 2.3 were employed to demonstrate the 2D and 3D molecular interactions of ligands **3**, **4**, **7** and **8** in the active sites of 6LU7 receptors. These interactions of **3**, **4**, **7**, **8** and ribavirin are presented in **Fig. 7**, while the affinity is listed in **Table 4**.

Table 4. The affinity of **3**, **4**, **7**, **8** and ribavirin docked in COVID-19 main protease (6LU7)

Product	Affinity (kcal/mol)
3	-4.2
4	-4.8
7	-6.5
8	-6.4
Ribavirine	-6.2

The binding energies (kcal/mol) for the 6LU7 crystal structure of the main protease of SARS-CoV-2 are in the order of $-6.5 > -6.4 > -6.2 > -4.8 > -4.2$ kcal/mol for product **7** > product **8** > ribavirin > product **4** > product **3** >, respectively. When compared to the standard antiviral drugs (ribavirin) and the other product under study, the product **7** > having triazoles and isoxazoles rings was found to have good antiviral activity, and we notice that its affinity is increased by the meta cyclization. The product **7** may therefore be the best antiviral medication, according to the results for the product **7** and ribavirin derivatives.

8. Conclusion

On the basis of B3LYP/6-31G(d,p), MEDT has used the concept to evaluate [3+2] cycloaddition reactions between **1** and **2** and **5** and **6**, with the conclusion that **2** and **6** present as electrophiles for this reaction, **1** and **5** will participate as nucleophiles. Furthermore, the regioselectivity observed experimentally is perfectly confirmed by Parr function analysis. Moreover, these reactions are irreversible, thanks to their highly exergonic nature. ELF analysis of the most likely reaction trajectory asserts that this action occurs according to an uncoordinated mechanism, also, a docking investigation of products studied docked in Covid-19 major protease (6LU7) in comparison to Ribavirin was done; the findings indicate compounds **7** could serve as an antiviral drug.

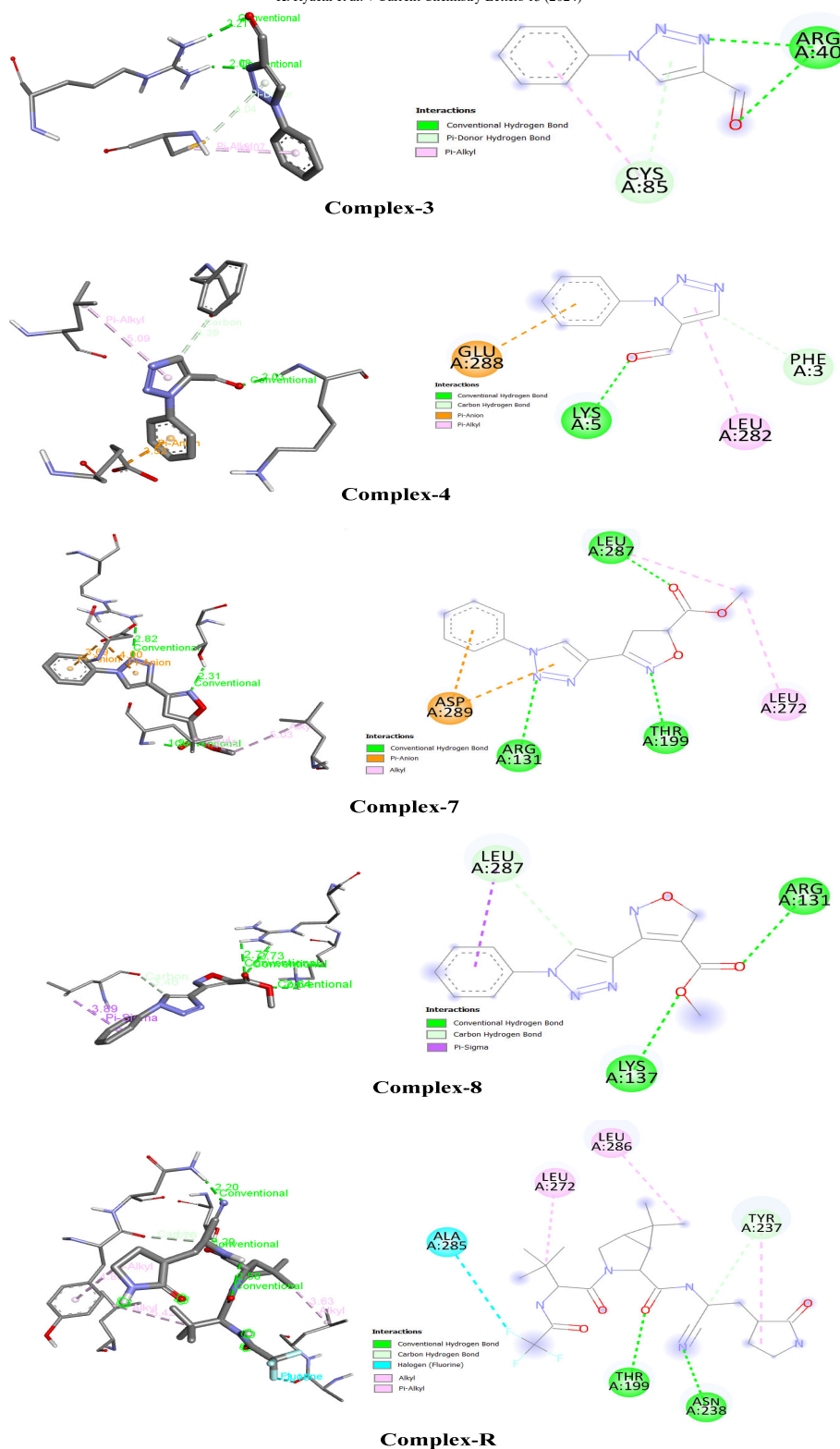


Fig. 7. 2-3-D-Visualizations of docking of 3, 4, 7, 8 docked in COVID-19 main protease (6LU7) compared to the Ribavirin.

Declaration of Competing Interest

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

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Ethics approval

The manuscript is prepared in compliance with the Ethics in Publishing Policy as described in the Guide for Authors.

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