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Properties and Reactivities of Niclosamide in Different Media, a Potential Antiviral to Treatment of COVID-19 by **Using DFT Calculations and Molecular Docking**

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Abstract: In this work, the structural, electronic, topological and vibrational properties of potential antiviral to treatment of COVID-19, niclosamide (NCL) have been studied in different media together with its reactivities by combination of DFT calculations with molecular docking. Properties of two most stable conformers of niclosamide (C1 and C2) were reported in gas phase and water, ethanol and chloroform solvents. Calculations using the integral equation formalism variant polarised continuum (IEFPCM) and solvation methods in the different solutions have revealed solvation energy values for C1 and C2 in aqueous solution (ΔG_C = -78.43 and -64.53 kJ/mol, respectively) comparable with that observed for the antiviral agent zalcitabine (-78.92 kJ/mol). Probably, the high stability of C1, predicted by NBO studies, explains the experimental existence of C1 in the solid phase. Comparisons of frontier orbitals with eleven antiviral agents have evidenced the high reactivity of C2 slightly higher than brincidofovir, an antiviral agent used in the treatment to *ebola* disease. Possibly, the presence of deactivating groups (NO₂ and Cl) in the chloro-4-nitrophenyl and hydroxybenzamide rings of both forms of NCL could explain the higher reactivities predicted in the different media. Here, the harmonic force fields and force constants for both forms are reported together with the assignments of 80 vibration modes expected in the experimental infrared spectrum of NCL. The predicted UV-Vis spectra in the different solvents suggest the presence of both forms of NCL in solution. Molecular docking results were discussed basing on the type of interaction between the ligands and several amino acid residues.

Keywords: Niclosamide; vibrational spectra; molecular structure; descriptor properties; DFT calculations

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1. Introduction

The great world problem at present is the treatment of the deadly coronavirus disease 2019 named "severe acute respiratory syndrome coronavirus 2" (SARS-CoV-2) that is affecting the world population [1-20]. The identification and characterization of novel 2019nCoV were performed by Zhu et al. [3] whose have revealed that the new 2019-nCoV presents from 75 to 80% equivalent to the SARS-CoV and even more closely related to several bat coronaviruses [4]. The crystal structure of 2019-nCoV main protease in complex with an inhibitor N3 was identified [5]. In contrast, the inhibitor N3 was previously published by Yang https://biointerfaceresearch.com/

et al. [6]. In this context, the investigations are directed to the development of therapeutic agents and vaccines for COVID-19 and related human coronavirus diseases [9,10]. The need to find an effective antiviral agent is urgent; so many researchers are working on the design of effective antiviral agents with few side effects. So, recently Xu et al. [12] have suggested niclosamide (NCL) as a potential antiviral agent because it drug is effective against various viral infections with nanomolar to micromolar potencies such as SARS-CoV, MERS-CoV, ZIKV, HCV, and human adenovirus. Structures of compounds clinically used as antiviral drugs for the treatment of numerous infections, such as human immunodeficiency virus (HIV), of chronic hepatitis B virus (HBV) and hepatitis C virus (HCV), of herpes simplex virus (HSV) and/or varicellazoster virus (VZV), or cytomegalovirus (CMV, of influenza virus and, of respiratory syncytial virus (RSV) have been carefully detailed by Clerk including their activity spectral, mechanisms of action, principal indications and administration [21,22]. Hence, those compounds together with other antiviral agents are known from a biological point of view, while the structural, electronic, topological, and vibrational properties only for some of them were studied combining available experimental results with theoretical DFT calculations [23-32]. Here, the structural, electronic, topological, and vibrational properties of two conformers most stable of niclosamide, 5-Chloro-N-(2-chloro-4-nitrophenyl)-2-hydroxybenzamide, were reported in the gas phase and water, ethanol and chloroform solvents by using DFT calculations and molecular docking. It drug was studied in different media because studies on cocrystals of NCL have evidenced its low solubility and high permeability [33] and, for these reasons, here, investigations of reactivities and behaviors in the different solvents for both forms were performed. Those properties, together with the NCL solubility in different solvents, are essential, taking into account the expected potential biological activities and the presence of donors and acceptors groups in the structure. Here, the solvation energy values, atomic charges, stabilization energies, molecular electrostatic potentials, bond orders, and reactivities were evaluated as a function of the solvent's permittivity values. On the other hand, the bands observed in the experimental available infrared spectrum were assigned combining the B3LYP/6-31G* calculations with the SQMFF methodology and the Molvib program [34-36]. Hence, the harmonic force constants in the different media are also reported. The predicted properties for NCL were compared with those published for other antiviral agents [23-32]. In addition, a molecular docking study was performed for NCL because it is an essential method in molecular modeling applications and drug discovery.

2. Materials and Methods

The initial structure of NCL was taken from the CIF file because its experimental structure was determined by Sanphui et al. [37] by using X-ray diffraction. The *GaussView* program and the Revision A.02 of the Gaussian program [38,39] were employed to optimize the structures of NCL in the different media with the functional hybrid B3LYP/6-31G* method [40,41]. That basis set was used because the properties of compared antiviral compounds were published with that level of theory. Both IEFPCM and universal solvation methods employed here to consider the solvent effects [42-44]. With the optimized structure in the gas phase, the potential energy surfaces (PES) were investigated for variations of dihedral N7-C11-C10-C14 and C10-C14-O3-H29 angles at the same level of theory. Then, two stable structures, named C1 and C2, were found in the PES, where C2 presents the global minimum and where both can be seen in Figure 1.



Figure 1. Molecular structures of the two most stable conformers of niclosamide (5-Chloro-N-(2-chloro-4nitrophenyl)-2-hydroxybenzamide), identifications of both rings and atoms numbering.

Here, it is necessary to clarify that the experimental structure of NCL determined by Sanphui et al. [37] corresponds to conformer C1, but the theoretically most stable structure of NCL is C2. Volumes and their differences were computed with the Moldraw program [45]. Versions 3.2 and 2000 of NBO and AIM programs respectively were used to calculate atomic charges, molecular electrostatic potentials, bond orders, and topological properties [46-48] while the reactivities and behaviors in the different media were predicted by using frontier orbitals and equations reported for some typical descriptors [49-56]. Usual internal coordinates of NCL, transferable scaling factors, and the scaled mechanical force field (SQMFF) methodology, together with version 7.0 of the Molvib program, were used to calculate the harmonic force fields and force constants [34-36]. Then, the vibrational study was performed using potential energy distribution (PED) contributions ≥ 10 %.

All molecular docking calculations for niclosamide (NCL) are made using iGEMDOCK [57] on the basis of GEMDOCK (Generic Evolutionary Method for Docking Molecules) scoring function [58]. The choice of ligands was mainly based on their antiinflammatory and antiviral activities. It is worthy of mentioning that the novel 2019-nCoV has the same symptoms as a flue. It can cause fatigue, fever, and cough, which may be accompanied by nasal congestion, runny nose, headache, expectoration, diarrhea, etc. The election was performed, taking into account that information. Before starting the molecular docking calculations, the structure of Coronavirus 2019 (COVID-19) is extracted from the Protein Data Bank [5] of the Structural Bioinformatics Research Laboratory (RCSB) (PDB ID: 6LU7). Subsequently, it is cleaned of possible undesirable entities such as water molecules by the Discovery Studio program [59]. NCL was tested as 2019-nCoV inhibitors and was docked at the binding site of 2019-nCoV to shed light on their potential binding modes. The p-tau protein-ligand complex was calculated by measuring various non-polar and polar interactions, for instance, van der Waal's interactions, electrostatic interactions, and H-bonding interaction interactions for each ligand.

3. Results and Discussion

3.1. Optimizations and properties in different media.

Both stable molecular structures of NCL and atoms numbering are shown in Figure 1, together with the identification of two rings. Rings containing both chloro and nitrophenyl groups are named R1, while R2 is named to the rings that contain to hydroxybenzamide moieties. Calculated total and corrected by zero-point vibrational energy (ZPVE), dipole moments (μ) and volumes (V) of both conformers of NCL in the gas phase and aqueous, ethanol and chloroform solutions by using B3LYP/6-31G* level of theory together with the

permittivity's (ε) values can be seen in Table 1. Higher energy values are observed for C1 and C2 in chloroform solution, although the dipole moment values have higher values in water, and its higher volumes are observed in ethanol. Besides, for both conformers, the dipole moment values increase according to increases the permittivity of the solvent, although the volume values do not follow a defined trend. Variations energy values among both forms show that C1 and C2 could exist in the different solutions because in the gas phase ΔE is 20.46 kJ/mol while the values decrease in chloroform, ethanol, and water respectively to 12.06, 6.82 and 7.34 kJ/mol. This way, the values predicted in solutions are low enough, as compared with the values in the gas phase. Table 2 are presented corrected solvation energies and uncorrected by ZPVE energies together with the volumes variations for C1 and C2 of NCL in the gas phase and the three solvents by using the B3LYP/6-31G* level of theory.

theory. Permittivity's (ϵ) values of all species are also included.								
	B3LYP/6-31G* Method							
	Conformer C1							
Species	E (Hartrees)	E (Hartrees) E ZPVE μ (D) V (Å ³)						
Gas	-1830.9136	-1830,7170	7,95	278,9	1			
Chloroform	-1830.9391	-1830,7426	9,77	278,8	4.7113			
Ethanol	-1830.9382	-1830.7423	10,93	285,2	24.852			
Water	-1830.9294	-1830,7337	11,25	282,8	78.3553			
	Conformer C2							
Species	E (Hartrees)	E ZPVE	μ (D)	V (Å ³)	3			
Gas	-1830.9214	-1830.7241	3.96	281.5	1			
Chloroform	-1830.9437	-1830.7465	4.74	286.3	4.7113			
Ethanol	-1830.9408	-1830.7440	5.43	287.7	24.852			
Water	-1830.9322	-1830.7356	5.65	286.3	78.3553			

Table 1. Calculated total and corrected by ZPVE energies (*E*), dipole moments (μ), and volumes (V) of niclosamide in the gas phase and aqueous, ethanol and chloroform solutions by using B3LYP/6-31G* level of

Table 2. Corrected solvation energies ($\Delta G_{C/ZPVE}$) and uncorrected by ZPVE energies (ΔG_C) and volumes variations (ΔV) of niclosamide in the gas phase and aqueous, ethanol and chloroform solutions by using B3L/YP/6-31G* level of theory.

B3LYP/6-31G* Method								
		Conformer (21					
Species	ΔG_{un}	ΔG_{un} ΔG_{ne} ΔG_C $\Delta V (Å^3)$ $\Delta G_{C/ZM}$						
Chloroform	-66.89	-21.53	-45.36	-0.1	-45.62			
Ethanol	-64.52	8.07	-72.59	6.3	-78.89			
Water	-41.44	36.99	-78.43	3.9	-80.79			
Conformer C2								
Species	ΔG_{un}	ΔG_{ne}	ΔG_{C}	$\Delta V (Å^3)$	$\Delta G_{C/ZPVE}$			
Chloroform	-58.49	-21.36	-37.13	4.8	-37.39			
Ethanol	-50.88	8.78	-59.66	6.2	-60.97			
Water	-28.33	36.20	-64.53	4.8	-66.36			

Here, the ($\Delta G_{C/ZPVE}$) value implies two corrections to solvation energy value, which are corrections by non-electrostatic terms and by ZPVE while with (ΔG_C) is identified to that value only corrected by non-electrostatic terms. The variations of dipole moment, volume, and solvation energy values as functions of permittivity's solvents for C1 and C2 are shown in Figure 2. From Table 1 and Figure 2a, it is observed for both forms the increase of μ with the permittivity of solvent where clearly the relationships between both properties are non-linear

and practically follow the same tendency, but the values are higher for C2. The magnitudes of dipole moment values and vectors for C1 are different from C2 in the three solutions, as compared with the corresponding values in the gas phase but practically are constants in the three solutions, as observed in Figure S1. The changes in the position of the OH group generate significant variation in the dipole moment vector of C2.



Figure 2. Variations of dipole moment (a) volume (b) and corrected solvation energy values for the two most stable conformers C1 (c) and C2 (d) of niclosamide as a function of permittivities of solvents.

Concerning volumes, Figure 2b shows that the V values for C1 and C2 increase with the permittivity from NCL in the gas phase up to NCL in ethanol solution but, later, the values decrease slightly in water. Note that the V values in C1 are higher than the corresponding to C2. The relationships between V and ε values are also non-linear. If now the predicted solvation energies (ΔG_C) are analyzed from Table 2 and Figures 2c-2d we observed that the values are higher for C1 and C2 in water ($\Delta G_C = -78.43$ and $\Delta G_C/ZPVE = -80.79$ for C1 and $\Delta G_C = -64.53$ kJ/mol, $\Delta G_C/ZPVE = -66.36$ kJ/mol for C2). Moreover, for both forms, the ($\Delta G_C/ZPVE$) and (ΔG_C) values decrease with ε , but the values are most negative corrected by ZPVE ($\Delta G_C/ZPVE$). Figures 2c-2d show few differences between both curves for C1 and C2 and where the solvation energies values in chloroform remaining practically constants. At the same time, a slight increase it is observed in ethanol and water (red curve, most negative values).

In this work, the solvation energy (Δ Gc) in aqueous solution were also considered in function of acceptors and donors groups present in both structures of niclosamide because these groups are essential to analyze its oral bioavailability and absorption, as suggested by Veber et al. [60] and Lipinski et al. [61]. Hence, the H bonds donors (N-H or OH groups) and acceptors (N and O atoms) present in C1 of NCL structure (because it has higher (Δ Gc) and, other atoms (2 Cl atoms) and groups (NO₂, 2 phenyl rings) were compared in Table 3 with solvation energy values reported for other ten antiviral agents in the same medium by using the same level of theory. Thus, the value for the conformer C1 of NCL in water is compared with the same method, isothiazol [24], zalcitabine [25], emtricitabine [28], trifluridine [26], thymidine

[23,29], idoxuridine [27], ribavirin [31], cidofovir and brincidofovir [32] and foscarnet [30]. The structures of all antiviral compounds can be clearly seen in Figure S2.

anu	and O atoms present in ten antivital species in aqueous solution by using the hybrid B5E 11/0-510 inteniod.									
N°	Species	ΔG_{C}	N-H	O-H	0	C=O	Ν	ToTal	Groups	Rings
1	Isothiazol ^{b,}	-37.51	1				2	3	SH, C≡N	R5,R6
2	Chloroquine ^a	-52.06	1				3	4	Cl	2 R6
3	Niclosamide ^a	-78.43	1	1	4	1	2	9	2 Cl, NO ₂	2 R6
4	Zalcitabine ^c	-78.92	2(NH ₂)	1	3	1	3	10		R5,R6
5	Emtricitabine ^d	-100.88	2(NH ₂)	1	3	1	3	10	F	R5,R6
6	Trifluridine ^e	-113.85	1	2	5	2	2	12	CF ₃	R5,R6
7	Thymidine ^f	-116.16	1	2	5	2	2	12	CH ₃	R5,R6
8	Idoxuridine ^{g,#}	-124.50	1	2	5	2	2	12	Ι	R5,R6
9	Ribavirin ^h	-141.85	2(NH ₂)	3	5	1	4	15		2R5
10	Cidofovir ⁱ	-169.21	2(NH ₂)	3	6	1	3	15	H_2PO_3	R6
11	Foscarnet ^j	-219.64		12	5	2		19	3 Na,	
12	Brincidofovir ⁱ	-227.34	2(NH ₂)	2	7	1	3	15	HPO ₃	R6

Table 3. Uncorrected solvation energies by ZPVE energies (ΔG_C) and numbers of N-H and O-H groups and N and O atoms present in ten antiviral species in aqueous solution by using the hybrid B3LYP/6-31G* method.

^aThis work, ^bFrom Ref [24], ^cFrom Ref [25], ^dFrom Ref [28], ^cFrom Ref [26], ^fFrom Ref [23,29], ^gFrom Ref [27], ^bFrom Ref [31], ⁱFrom Ref [32], ^jFrom Ref [30]. [#]Idoxuridine calculated by using B3LYP/3-21G* calculations.

Note that in the isothiazol structure, 3-mercapto-5-phenyl-4-isothiazolecarbonitrile, there is an S-H instead of N-H or OH as the H bond donor whiles the structures of acid, its trisodic anhydrous and hexahydrated salts are observed for foscarnet. For idoxuridine, the ΔG_C values were calculated by using B3LYP/3-21G* (-124.50 kJ/mol) and B3LYP/Lanl2dz (-162.76 kJ/mol) calculations. Here, the ΔG_C value for idoxuridine (-124.50 kJ/mol) with the B3LYP/3-21G* method was considered because it presents a higher energy value (most negative). Note that all compared species have rings in their structures, with the exception of foscarnet because it is a trisodic salt hexahydrated and, for this reason, it drug presents a higher total number of acceptor/donor groups. In foscarnet, each water molecule was also considered as two donor O-H groups. The trisodic salt, foscarnet, has the higher ΔG_C value probably due to B3LYP/6- $311++G^{**}$ method and/or to its cationic form present in aqueous solution. In Figure 3 can be seen the quick increase of ΔG_C with the total number of acceptor/donor groups from isothiazol up to foscarnet and then decrease up to brincidofovir. The properties of cidofovir to brincidofovir were previously published in the gas phase [32], but the solvation energy value of brincidofovir in water is here reported for the first time. Both forms of NCL presents only total 9 groups, as compared with the other antiviral agents where we can see that trifluridine [26], thymidine [23,29] and idoxuridine [27] have the same total number of acceptor/donor groups because trifluridine is different from thymidine in a CF₃ group while idoxuridine has a iodide atom. Moreover, these three antiviral agents have the same five and six member's rings in their structures. Here, C1 and C2 in aqueous solution have solvation energy values of ΔG_{C} = -78.43 and -64.53 kJ/mol, respectively between those observed for isothiazol (-37.51 kJ/mol) and zalcitabine (-78.92 kJ/mol). These results show that despite less negative solvation energy values predicted for C1 and C2 in aqueous solution, as compared with other antiviral agents, the presence of deactivating groups (NO₂ and Cl) in the chloro-4-nitrophenyl and hydroxybenzamide rings could explain the higher reactivities observed for NCL in different media.



Figure 3. Total number of acceptors and donors groups of eleven antiviral agents as a function of corrected solvation energy values compared with the corresponding to C1 by using B3LYP/6-31G* level of theory.

3.2. Geometrical parameters in different media.

The geometrical parameters calculated for both conformers of NCL in the gas phase and chloroform, ethanol, and water by using the functional hybrid B3LYP with the 6-31G* basis set were compared with the corresponding experimental ones determined for C1 of NCL by Sanphui et al. [37] by X-ray diffraction. Comparisons between both experimental and theoretical parameters were carried out with the root-mean-square deviations (RMSD), which are presented in Table 4 for C1, while in Table S1 are summarized for C2. Analyzing deeply the values for both conformers, a better correlation is observed in bond lengths of C1, as expected because the experimental structure reported for NCL by Sanphui et al. [37] corresponds to C1. Hence, the bond lengths values in C1 are between 0.025 and 0.022 Å while in C2 between 0.026 and 0.024 Å. The differences in the bond angles in C1 are between 3 and 2.9 ° while in C2, the values are slightly lower (1.9 and 1.8 °). Both forms show higher variations in the dihedral angles. Thus, the dihedral Cl2-C20-C16-C10 angle for C1 in the gas phase has a positive value while in the three solutions, change the signs to negative. On the contrary, that dihedral angle in C2 presents in all media positive values. The dihedral O6-N8-C15-C18 angle in C1 shows negative values in chloroform and ethanol solutions, while this angle presents a negative sign only for C2 in chloroform solution.

The dihedral O4-C11-C10-C14 angles for C1 in ethanol and water have negative values, while for C2 in all media, the angles show positive values. These changes in the dihedral angles for C1 and C2 in the three solutions evidence that better correlations are obtained in the gas phase (2.4-2.1 °) because the presence of acceptors and donor groups H bonds suggest that these groups are solvated. Therefore, both optimized C1 and C2 structures can be used to perform vibrational studies.

Parameters	Gas	Chloroform	Ethanol	Water	Exp ^b
	1	Bond lengt	hs (Å)		1
C11-O4	1.225	1.228	1.231	1.233	
C14-O3	1.370	1.363	1.361	1.364	1.331
N7-C9	1.394	1.391	1.389	1.390	1.395
N7-C11	1.383	1.383	1.381	1.380	1.361
C10-C11	1.510	1.505	1.502	1.500	1.540
C12-Cl1	1.761	1.761	1.760	1.759	1.730
C20-Cl2	1.755	1.763	1.765	1.764	1.737
C15-N8	1.464	1.456	1.451	1.449	1.462
N8-O5	1.231	1.234	1.236	1.237	1.220
N8-O6	1.232	1.234	1.236	1.237	1.219
RMSD	0.022	0.023	0.024	0.025	
	1	Bond ang	les (°)	1	1
N7-C11-O4	123.1	122.8	122.6	122.5	
N7-C11-C10	116.9	116.9	116.9	117.1	
C10-C11-O4	119.9	120.1	120.3	120.2	
C10-C14-O3	119.0	119.0	118.9	119.0	
C19-C14-O3	120.5	120.5	120.8	120.7	111.1
C9-N7-C11	127.6	127.9	128.1	128.1	129.1
C15-N8-O5	117.6	118.0	118.3	118.4	118.1
C15-N8-O6	117.7	118.2	118.5	118.6	118.5
O6-N8-O5	124.6	123.7	123.0	122.9	123.2
N8-C15-C17	118.8	118.8	118.9	118.8	118.2
N8-C15-C18	119.8	119.7	119.7	119.7	119.3
Cl1-C12-C9	120.2	120.1	120.1	120.1	119.3
Cl1-Cl2-Cl7	117.8	117.7	117.7	117.7	118.0
Cl2-C20-C16	119.8	119.5	119.5	119.5	119.5
Cl2-C20-C21	119.5	119.5	119.4	119.4	119.3
RMSD	2.9	2.9	3.0	2.9	
	1	Dihedral an	gles (°)		1
C9-N7-C11-O4	-0.0	-179.9	-0.9	-0.7	
C9-N7-C11-C10	179.9	-0.0	178.0	178.7	
N7-C9-C12-Cl1	0.0	0.0	0.2	-0.2	-3.6
O6-N8-C15-C18	179.9	-179.9	-179.8	179.7	179.9
N7-Cl1- C10-C14	-0.0	-0.0	6.6	4.0	
C11-C10-C14-O3	0.0	-0.0	1.0	0.4	
Cl2-C20-C16-C10	180.0	-180.0	-179.9	-179.8	177.8
O4-C11-C10-C14	179.9	179.9	-174.3	-176.3	
RMSD	2.4	293.0	292.9	206.5	

 Table 4. Comparison of calculated geometrical parameters for the conformer C1 of niclosamide in gas and chloroform, ethanol and water solutions compared with the corresponding experimental ones in the solid phase.

 B3LYP/6-31G*a

^aThis work, ^bRef [37].

3.3. Atomic MK and Mulliken charge, molecular electrostatic potential, and bond orders.

Atomic charges and the molecular electrostatic potentials in a molecule are useful parameters to describe and explain the different reaction sites and behaviors of species in diverse mediums. In addition to acceptors and donor groups, the presence of two deactivating groups (NO₂ and Cl) in the chloro-4-nitrophenyl ring and of Cl atoms in both rings of C1 and C2 could easily justify the high reactivities predicted in the different media, as we will see later. Hence, atomic Merz-Kollman (MK) and Mulliken charges, molecular electrostatic potentials (MEP), and bond orders, expressed as Wiberg indexes for C1 and C2 in different media by using B3LYP/6-31G* level of theory are presented in Table S2. The behaviors of MK and Mulliken charges on the eight Cl, O, and N atoms belonging to C1 and C2 in the four media can be seen in Figure S3. The analysis of parameters shows practically the same behaviors of MK and Mulliken charges in the four media and where the MK charges show positive values on all atoms with the exception of N8. On the contrary, Mulliken charges evidence positive values on Cl1 and N8 atoms of C1 and C2 but with null values on Cl1. The lower MK and Mulliken charge values for C1 and C2 are observed on the O3 and N7 atoms, probably due to that these atoms are involved in the H bonds formations. In C1, the two C11-O4---H23 and N7-H22...O3 bonds are formed while, in C2, four H bonds are formed (C11-O4...H23, O3-H29...O4, C12-Cl1...H22 and N7-H22...H24 bonds). Therefore, C2 is most stable than C1, as was previously analyzed.

Regarding the molecular electrostatic potentials, the values on the eight considered atoms are approximately the same in the four media, as observed in Table S2. However, when the mapped MEP surfaces are graphed in Figure S4, different colorations for both forms are found. Thus, in C2 strong red colors are observed on the O atoms of NO₂, C=O, and OH groups while the colorations on the NO₂ and C=O groups are weak in C1. This fact could be explained taking into account the four H bonds in C2, for which, the R1 ring is less deactivate because the C11 atom is forming the C12-C11...H22 bond while in C1 the C11 atoms are not involved in a H bond. On the other hand, the strong blue color in C2 is observed on the N7-H22 bond while in C1 is observed on the O3-H29 bond and in this form that NH bond is involved in the formation of N7-H22...O3 bond. Then, the nucleophilic and electrophilic sites can be identified respectively with red and blue colors, while neutral or inert regions are identified by green colors. Here, the strong red and blue colorations observed for C2 could be associated with the higher reactivity predicted for this form, as compared with the C1 form.

In addition, for C1 and C2, bond orders (BO) expressed as Wiberg indexes were also calculated due to the presence of deactivating and acceptors and donors groups in its structures. Accordingly, in Table S2, the BO of those eight atoms of C1 and C2 calculated in the four media by using the B3LYP/6-31G* level of theory are presented. The careful analysis of data shows that the BO values for the different atoms follow the tendency N > O > CI. Thus, the Cl and O atoms have approximately the same values in both forms and in the four media, but the N8 atoms have higher values than those observed for the N7 atoms, as expected because the N8 atoms belong to the NO₂ groups and they are linked to O atoms. Here, the BO values do not any show significant differences between both conformers in the four media.

3.4. NBO and AIM studies.

Due to the presence of acceptors and donors groups, of two deactivating groups (NO₂ and Cl) in the chloro-4-nitrophenyl ring and Cl atoms in both rings of C1 and C2, calculations https://biointerfaceresearch.com/

of Second-Order Perturbation Theory Analysis of Fock Matrix in NBO Basis by using NBO analysis and of topological properties by using Bader's theory of atoms in molecules (AIM) are valuable tools necessaries to investigate the existence of different interaction's types, such as H bonds or inter and intra-molecular interactions [45-48]. Hence, acceptor-donors energies and topological properties were predicted for C1 and C2 of NCL in the four media by using the versions 3.1 and 2000 of NBO and AIM programs, respectively [46,47]. Thus, the main delocalization energies for C1 and C2 in the gas phase, chloroform, ethanol, and water solutions by using the B3LYP with the 6-31G* basis set are presented in Tables S3 and S4, respectively. The exhaustive analysis of results show that C1 presents five different interactions ($\Delta E_{\pi \to \pi^*}$, $\Delta E_{\pi \to n}$, $\Delta E_{n \to \pi^*}$, $\Delta E_{n \to \sigma^*}$ and $\Delta E_{\pi^* \to \pi^*}$) while for C2 in the different media are observed seven interactions ($\Delta E_{\pi \to \pi^*}$, $\Delta E_{\pi \to n^*}$, $\Delta E_{\pi \to \pi^*}$, $\Delta E_{n \to \pi^*}$, $\Delta E_{n \to \sigma^*}$, $\Delta E_{n \to n^*}$ and $\Delta E_{\pi^* \to \pi^*}$). On the other hand, due to the different positions of O3-H29 group in C1 and C2 two new $\Delta E_{\pi \to n^*}$ and $\Delta E_{n \to n^*}$ interactions are observed in C2 different from C1. Besides, the $\Delta E_{\pi \to \pi^*}$ and $\Delta E_{\pi^* \to \pi^*}$ transitions in C1 present higher values than those observed in C2 while, on the contrary, the predicted $\Delta E_{n \to \pi^*}$ transitions in C2 is higher than C1. A very important resulted is observed in the same values of $\Delta E_{n\to\sigma^*}$ transitions related to the lone pairs of O4, O5, and O6 atoms of both conformers (504.67/504.59 kJ/mol). Probably, such observation can be explained because the positions of these atoms in chloro-4-nitrophenyl rings and C=O groups in both conformers present the same positions in both forms. In contrast, the other hydroxybenzamide rings are different in C1 and C2. This NBO study clearly shows the higher stability of C1 than C2.

According to Bader's theory of atoms in molecules (AIM), the topological properties were computed for C1 and C2 in the four media by using the AIM2000 program in order to investigate H bonds or inter and intra-molecular interactions [47,48]. Thus, the electron density, $\rho(r)$, the Laplacian values, $\nabla^2 \rho(r)$, the eigenvalues ($\lambda 1$, $\lambda 2$, $\lambda 3$) of the Hessian matrix and the $\lambda 1/\lambda 3$ ratios were calculated for C1 and C2 in the bond critical points (BCPs) and the ring critical points (RCPs) by using the B3LYP/6-31G* method. Here, it is necessary to remember that the H bond interactions present a ratio of $\lambda l/\lambda 3 < 1$ and $\nabla^2 \rho(r) > 0$. Those topological properties for C1 and C2 in the different media can be seen in Tables S5 and S6 together with the distances between the involved atoms. In C1, three new BCPs are predicted, which are the C11-O4…H23, C11-O4…H23 and C12-Cl1…H22 interactions and, for these reasons, three new RCPs appear (RCPN1, RCPN2 and RCPN3) while in C2, four new interactions (C11-O4-H23, C11-O4-H29, C12-C11-H22 and N7-H22-H24 interactions) and four new RCPs are observed (RCPN1, RCPN2 and RCPN3). Besides, in both forms are present the RCPs of both chloro-4-nitrophenyl and hydroxybenzamide rings (RCP1 and RCP2). All interactions of conformers C1 and C2 in the gas phase showing the geometry of all their BCPs and RCPs can be easily seen in the molecular graphics presented in Figure S5. These AIM studies show the high theoretical stability of C2.

3.5. Frontier orbitals and global descriptors studies.

The frontier orbital studies are very important in C1 and C2 of NCL because a recent study performed by Xu et al. [12] suggest to NCL as a potential antiviral agent due to its effectivity against various viral infections such as SARS-CoV, MERS-CoV, ZIKV, HCV, and human adenovirus. Hence, the reactivities of both forms are here predicted by using the differences between the HOMO and LUMO named gap, as recommended by Parr and Pearson [49]. Then, with the gap values the chemical potential (χ), electronegativity (μ), hardness (η),

softness (*S*), global electrophilicity index (ω) and global nucleophilicity index (*E*) descriptors are also calculated with known equations [50-56]. Then, all these properties for C1 and C2 in the gas phase, chloroform, ethanol, and aqueous solutions by using the hybrid B3LYP level of theory are presented in Table 5 together with the corresponding equations. C1 shows higher values in all media, as compared with C2, while slight reductions in gap values are observed for both forms in the different media when increasing the permittivities of solvents. Later, both conformations evidence higher reactivities in water.

nıc	niclosamide by using the hybrid B3LYP level of theory.						
B3LYP/6-31G*							
		Conformer	C1				
Orbitals	Gas	Chloroform	Ethanol	Water			
HOMO	-6.7457	-6.7484	-6.7511	-6.7566			
LUMO	-2.5252	-2.5524	-2.5715	-2.5824			
GAP	4.2205	4.1960	4.1796	4.1742			
		DESCRIPT	ORS				
χ	-2.1103	-2.0980	-2.0898	-2.0871			
μ	-4.6355	-4.6504	-4.6613	-4.6695			
η	2.1103	2.0980	2.0898	2.0871			
S	0.2369	0.2383	0.2393	0.2396			
ω	5.0912	5.1540	5.1985	5.2236			
Е	-9.7820	-9.7565	-9.7412	-9.7457			
		Conformer	C2				
Orbitals	Gas	Chloroform	Ethanol	Water			
HOMO	-6.5471	-6.5552	-6.5688	-6.5797			
LUMO	-2.7864	-2.8218	-2.8436	-2.8572			
GAP	3.7607	3.7334	3.7252	3.7225			
		DESCRIPT	ORS				
χ	-1.8804	-1.8667	-1.8626	-1.8613			
μ	-4.6668	-4.6885	-4.7062	-4.7185			
η	1.8804	1.8667	1.8626	1.8613			
S	0.2659	0.2679	0.2684	0.2686			
ω	5.7911	5.8879	5.9455	5.9809			
Е	-8.7751	-8.7520	-8.7658	-8.7822			

Table 5. Calculated chemical potential (μ), electronegativity (χ), global hardness (η), global softness (S), global
electrophilicity index (ω) and global nucleophilicity index (E) for conformers C1 and C2 of antiviral

 $\chi = - [E(LUMO) - E(HOMO)]/2; \mu = [E(LUMO) + E(HOMO)]/2; \eta = [E(LUMO) - E(HOMO)]/2; S = \frac{1}{2}\eta; \omega = \frac{\mu^2}{2}\eta; E = \mu * \eta$

When the global electrophilicity index (ω) and global nucleophilicity index (*E*) are analyzed we observed that the low reactivities that evidence both forms could be attributed to the high values of (ω) and most negative values of (*E*). Then, these properties for C1 and C2 of NCL are compared with those reported for antiviral chloroquine, isothiazol [24], zalcitabine [25], emtricitabine [28], trifluridine [26], thymidine [23,29], idoxuridine [27], ribavirin [31], cidofovir and brincidofovir [32] and foscarnet agents [30] in Table 6 at different levels of theory. When the values are deeply analyzed, quickly it is observed that C2 in all media presents slightly lower values (3.7601-3.7225 eV) and higher reactivity than brincidofovir (3.7715 eV), the antiviral agent used in the treatment to *Ebola* disease. In comparison, the values of C1 are slightly higher (4.2205-4.1742 eV). Hence, C2 of NCL is the most reactive antiviral agent than those eleven compared antiviral ones.

	B3LYP Method					
Frontier orbitals		6-3	31G*		3-21G*	6-31G*
(6 v)	Chloroquine ^a	Isothiazol ^b	Zalcitabine ^c	Emtricitabine ^d	Trifluridine ^e	Thymidine ^f
HOMO	-5.5708	-6.692	-6.1138	-6.1825	-6.8882	-6.1061
LUMO	-1.1137	-2.185	-0.7543	-1.0629	-1.3006	-0.6313
GAP	4.4571	4.507	5.3595	4.9336	5.5876	5.4748
]	DESCRIPTORS	5		
χ	-2.2286	-2.2535	-2.6798	-2.5598	-2.7938	-2.7374
μ	-3.3423	-4.4385	-3.4341	-3.6227	-4.0944	-3.3687
η	2.2286	2.2535	2.6798	2.5598	2.7938	2.7374
S	0.2244	0.2219	0.1866	0.1953	0.1790	0.1827
ω	2.5063	4.3710	2.2003	2.5635	3.0002	2.0728
Е	-7.4484	-10.0022	-9.2024	-9.2734	-11.4389	-9.2215
			B3LYP Method	1		
		6-31G*		6-311++G**	6-31G*	
Frontier orbitals	Idoxuridine ^{g,#}	Ribavirin ^h	Cidofovir ⁱ	Foscarnet ^j	Brincidofovir ⁱ	
HOMO	-6.2600	-6.7958	-5.9366	-6.9135	-5.5435	
LUMO	-1.2438	-0.9530	-0.6401	-0.6413	-1.772	
GAP	5.0162	5.8428	5.2965	6.2722	3.7715	
]	DESCRIPTORS	5		
χ	-2.5081	-2.9214	-2.6483	-3.1361	-1.8858	
μ	-3.7519	-3.8744	-3.2884	-3.7774	-3.6578	
η	2.5081	2.9214	2.6483	3.1361	1.8858	
S	0.1994	0.1712	0.1888	0.1594	0.2651	
ω	2.8063	2.5691	2.0416	2.2749	3.5474	
E	-9.4101	-11.3187	-8.7084	-11.8463	-6.8976	

Table 6. Frontier molecular HOMO and LUMO orbitals, gap and chemical potential (μ), electronegativity (χ), global
hardness (η), global softness (S), global electrophilicity index (ω) and global nucleophilicity index (E) for antiviral agents
calculated at different levels of theory by using the functional hybrid B3LYP

^aThis work, ^bFrom Ref [24], ^cFrom Ref [25], ^dFrom Ref [28], ^eFrom Ref [26], ^fFrom Ref [23,29], ^gFrom Ref [27], ^hFrom Ref [31], ⁱFrom Ref [32], ^jFrom Ref [30], [#]Idoxuridine calculated by using B3LYP/3-21G* calculations. $\chi = -$ [E(LUMO)-E(HOMO)]/2; $\mu =$ [E(LUMO) + E(HOMO)]/2; $\eta =$ [E(LUMO) - E(HOMO)]/2; $S = \frac{1}{2}\eta$; $\omega = \frac{\mu^2}{2}\eta$; $E = \mu * \eta$



Conformer C1 Conformer C2 4000 3000 2000 1000

Figure 5. Predicted Raman spectrum for C1 and C2

nitrophenyl)-2-hydroxybenzamide) in the gas phase

(5-Chloro-N-(2-chloro-4-

Figure 4. Experimental ATR spectrum of niclosamide (5-Chloro-N-(2-chloro-4-nitrophenyl)-2-

hydroxybenzamide) in mull of Nujol compared with the corresponding predicted for C1 and C2 in the gas phase by using $B3LYP/6-31G^*$ level of theory.

3.6. Vibrational study.

ul study.

niclosamide

by using B3LYP/6-31G* level of theory.

These vibrational studies are essential to identify both forms of NCL in all media by using the infrared and Raman spectra. The hybrid B3LYP/6-31G* calculations have optimized

of

the structures of C1 and C2 in all media with C_1 symmetries and only, 81 normal vibration modes with activities in both spectra are expected for both forms. Figure 4 shows the experimental available ATR spectrum of niclosamide in mull Nujol was taken from Ref [62] compared with that corresponding predicted for both forms in the gas phase by using the B3LYP/6-31G* method. Figure 5 are presented the predicted Raman spectra for both forms in the gas phase by using the same method.

For a better correlation, the theoretical Raman spectra in activities were transformed into intensities by using the corrections suggested in the literature [63,64]. Note that the group of IR bands intense and media intensities between 2954 and 2853 cm⁻¹ and the strong IR band at 751 cm⁻¹ are clearly assigned to nujol because it is hydrocarbon oil that absorbs in these regions.

The intense band predicted in the IR spectrum of C2 in the higher wavenumbers region, not observed in the experimental one, confirm that the form C1 is present in the solid phase, as experimentally was determined by Sanphui et al. [37] by using X-ray diffraction. The difference observed between experimental and theoretical spectra can be attributed to the calculations because these were performed in the gas phase, where the packing forces were not considered. Here, the harmonic force fields and the corresponding vibrational assignments for both forms of NCL were computed by using the SQMFF methodology, the normal internal coordinates, transferable scaling factors, and the Molvib program [34-36]. The experimental and calculated wavenumbers for both conformers C1 and C2 of NCL in gas phase are summarized in Table 7 by using the functional hybrid B3LYP with the 6-31G* basis set. Discussions only for some groups are presented below for the most important regions.

3.6.1. Band Assignments.

4000-2000 cm⁻¹ Region. In this region, for C1 and C2 are expected the six C-H and the N-H and O-H stretching modes corresponding to chloro-4-nitrophenyl and hydroxybenzamide rings, which are predicted between 3605 and 3047 cm⁻¹. Hence, the weak IR bands at 3581 and 3498 cm⁻¹ are assigned to the O-H and N-H stretching modes, respectively [23-32,50-56]. Then, the six C-H stretching modes are associated with the IR bands located between 3197 and 3079 cm⁻¹.

Experimental ^d	B3LYP/6-31G* Method ^a					
	C1			C2		
ATR ^c	SQM ^b	Assignments	SQM ^b	Assignments		
3581w	3605	vO3-H29				
3498w	3390	vN7-H22	3459	vN7-H22		
			3222	vO3-H29		
3197w	3154	vC13-H23	3153	vC13-H23		
3122sh	3128	vC17-H25	3128	vC17-H25		
3122sh	3119	vC18-H26	3121	vC18-H26		
3097m	3114	vC16-H24	3098	vC19-H27		
3079w	3096	vC19-H27	3086	vC21-H28		
3079w	3047	vC21-H28	3076	vC16-H24		
1686m	1707	vC11=O4	1672	vC11=O4		
1657m	1619	ρN7-H22	1615	νaNO ₂ , ρN7-H22		
1632m	1600	vC14-C19,vC20-C16	1606	vC20-C16		
1607s	1591	vaNO ₂	1588	vC13-C18		
1582s	1589	vC13-C18	1582	vaNO2, oN7-H22		

 Table 7. Observed and calculated wavenumbers (cm⁻¹) and assignments for conformers C1 and C2 of antiviral niclosamide in the gas phase by using the hybrid B3LYP level of theory.

Experimental ^d	B3LYP/6-31G* Method ^a				
1		C1		C2	
ATR ^c	SQM ^b	Assignments	SQM ^b	Assignments	
1567s	1581	vC10-C14,vC19-C21	1568	vC10-C14,vC19-C21	
1514s	1535	vC15-C18	1533	vC15-C18	
1470m	1482	βC16-H24	1480	βC19-H27	
1424m	1468	vC15-C17,vC9-C12	1462	vC15-C17	
1381w	1403	vC10-C16	1414	vC10-C16,vC20-C16	
1352s	1399	vC12-C17	1399	vC12-C17	
1331vs	1346	vsNO ₂	1366	δΟΗ	
1331vs	1316	vC9-C13	1348	vsNO ₂	
1331vs	1310	vC20-C16	1323	vC9-C13	
1295m	1282	vC14-O3	1312	vC14-O3	
1295m	1280	vC9-N7	1290	vC14-O3	
1231m	1251	βС17-H25,βС13-H23	1276	vC9-N7,βC13-H23,vC9-C12	
1231m	1234	βС17-Н25	1250	νC10-C16,βC16-H24,νC10-C11	
1198m	1210	vC10-C11	1235	βC17-H25	
1144w	1156	βС19-Н27,βС21-Н28	1199	vC14-C19	
1130w	1137	βC18-H26	1139	βC18-H26	
1116w	1114	δOH,vC11-N7	1133	βC21-H28	
1112w	1110	vC15-N8	1109	vC15-N8	
	1093	$\beta R_1(A1)$	1092	βR1(A1),vC20-C21	
1051w	1077	vC20-C21	1073	vC11-N7	
	1038	$\beta R_1(A2)$	1038	$\beta R_1(A2)$	
	992	γC18-H26	988	γC18-H26	
	949	γC16-H24	963	γC21-H28	
919w	924	γC21-H28	917	γC17-H25	
919w	916	γC17-H25	914	δC9N7C11	
908m	911	δC9N7C11, ρC11=O4	886	δNO ₂ ,vC15-N8	
908m	885	δNO ₂	863	γC16-H24	
	862	γС13-Н23	859	γС13-H23	
833m	842	$\beta R_3(A1)$	850	γC16-H24	
826m	815	δNO ₂ , ρC11=O4	837	γC19-H27	
826m	802	γС19-Н27	802	δNO ₂	
768sh	761	γC11=O4,γC10-C11	759	γC11=O4	
751s	743	γNO_2	742	γNO_2	
740sh	739	βR ₃ (A2)	730	βR ₃ (A2)	
711w	706	$\tau R_1(A2), \gamma C9-N7$	717	ρC11=O4,vC12-Cl1	
689w	682	γN7-H22,γC14-O3	698	$\tau R_1(A2)$	
	681	$\beta R_2(A2), \beta R_3(A1)$	690	тОН	
661w	661	$\tau R_1(A1)$	671	$\tau R_1(A1), \gamma C14-O3$	
(20)	653	$\beta R_2(A1)$	653	$\beta R_2(A2)$	
639w	637	$\beta R_2(A2)$	649	$\beta R_2(A1)$	
589w	560	βC14-O3	602	γN7-H22	
55/w	549	ρNO_2	557	ρΝΟ2	
541w	542	$\tau R_2(A1), \tau R_1(A2)$	535	γC15-N8	
532w	531	$\tau R_2(A1), \tau R_3(A1)$	525	$\tau R_2(A1)$	
504w			519	βC14-O3	
40/W	AEA	0.C0 N7	408	pC9-N/,VC12-C11	
453W	454	βC9-N7	447	pC11=04	
453W	446	$\tau K_3(A2), \gamma C12$ -Cll	442		
435W	435	τK ₃ (A1)	442	γC12-C11	
41/W	420	pC14-O3 6N/C11C10	428	γC10-C11	
400W	146	$v \subset 10 - C (11, \beta K_2(A1))$	403	ркз(A1)	
	251	vC20-C12	262		
	242	VC12-C11	244	VC20-C12	
	342	τκ3(Α1),γC20-Cl2	344	βC12-C11, ρNO ₂	

Experimental ^d	B3LYP/6-31G* Method ^a					
-	C1			C2		
ATR ^c	SQM ^b	Assignments	SQM ^b	Assignments		
	338	βC10-C11	335	τR ₃ (A1),γC20-Cl2		
	326	γC20-Cl2	321	γC20-Cl2		
	269	τОН	269	βC20-Cl2, δN7C11C10		
	265	βC20-Cl2				
	200	βC12-Cl1	208	βC15-N8		
			178	βC12-Cl1		
	174	$\tau R_2(A2)$	173	$\tau R_2(A2) \tau R_3(A2)$		
	165	γC15-N8				
	163	βC15-N8	164	γC15-N8,τN7-C11		
	148	βC20-Cl2,βC10-C11				
			132	τR ₂ (A1), τOH		
	127	$\tau R_2(A2), \tau R_3(A1)$	130	$\tau R_3(A1)$		
	107	$\tau R_2(A1), \tau R_3(A1)$	116	$\tau R_2(A2)$		
	64	δN7C11C10	67	τC11-C10,τN7-C9		
	59	τwNO ₂	53	τwNO ₂		
	48	τN7-C9	48	τC11-C10		
	23	τN7-C11	26	τN7-C11		
	11	τC11-C10	13	τN7-C9		

Abbreviations: v, stretching; δ , deformation in the plane; γ , deformation out of the plane; wag, wagging; τ , torsion; β_R , deformation ring τ_R , torsion ring; ρ , rocking; τw , twisting; δ , deformation; a, antisymmetric; s, symmetric; (A₁), Ring R1; (A₂), Ring R2. ^aThis work, ^bFrom scaled quantum mechanics force field; ^cFrom Ref [62].

2000-1000 cm⁻¹ Region. The C=O stretching mode is predicted in C1 at 1707 cm⁻¹ while in C2 at 1672 cm⁻¹. This difference in C2 is related to the two H bonds formed by the C=O group, as evidenced by the AIM study. Note that the intensities of these bands in both C1 and C2 conformers are different from that intense expected for these groups because the experimental ATR spectrum was recorded in mull of nujol, whose intense band at 751 cm⁻¹ overlaps with the band corresponding to NO₂ groups in this region diminishing the intensities ratio between these bands. The anti-symmetrical (v_a) and symmetrical (v_s) NO₂ stretching modes in C1 are assigned to the strong bands at 1607 and 1331 cm⁻¹ while for C2, the bands at 1657 and 1331 cm⁻¹ can be assigned to those two stretching modes. The C=C stretching modes of both rings with double and partial bond characters of two conformers can be assigned to the IR bands between 1632 and 1331 cm⁻¹ while the bending or in-plane deformation of CH groups (β C-H) can be attributed to the bands between 1470 and 116 cm⁻¹. The OH deformation modes are predicted for C1 and C2 in different regions. Thus these modes can be assigned to the bands at 116 and 1331 cm⁻¹, respectively. In C1, that mode is observed at lower wavenumbers due to the H bond formed between the N7-H22 and O3-H29 groups, as revealed by AIM analysis. The N-C stretching modes can also be assigned in this region because the SQM calculations predicted these modes in these regions [24-29,31,32].

1000-10 cm⁻¹ Region. In this region are expected the NO₂ deformation (δ), rocking (ρ) and out-of-plane deformation modes (γ) for C1 and C2. The SQM calculations predicted in approximately the same regions for both forms. Thus, the IR bands at 908, 826, and 751 cm⁻¹ can be assigned to those vibration modes, respectively. The three deformations and torsion ring modes of chloro-4-nitrophenyl, and hydroxybenzamide rings are assigned in those regions predicted by calculations, as detailed in Table 7 [23-32,50-56]. The remaining vibration modes are assigned as predicted by the SQM calculations and, in accordance with assignments reported for species with similar rings [23-32,50-56].

3.7. Force Fields.

The harmonic force constants were calculated for both forms of niclosamide because these parameters can predict the forces of different bonds and, hence, the formations of H bonds. Thus, the force constants for C1 and C2 were obtained from corresponding force fields by using the SQMFF methodology and the Molvib program [34-36]. Therefore, the harmonic force constants calculated for both forms of niclosamide in the gas phase with functional hybrid B3LYP and the 6-31G* basis set are given in Table 8 compared with those published for 2-(4 nitrophenyl)-4H-3,1-benzoxacin-4-one) at the same level of theory [65]. Some important differences can be observed when the values for C1 and C2 are compared. Hence, the differences in the values of f(vN-H), f(vO-H), f(vC=O), and $f(\delta O-H)$ force constants are related to the groups involved in the H bonds formation.

Force constant	Niclosamide ^a		2-(4 nitrophenyl)-4H-
	C1	C2	3,1-benzoxacın-4-one)
f(vN-H)	6.37	6.65	
f(vO-H)	7.27	5.77	
f(vC=O)	11.29	10.31	12.758
$f(vNO_2)$	9.64	9.69	9.735
$f(vN-C)_{NH}$	6.02	5.99	7.590
$f(vC-N)_{NO2}$	4.15	4.12	
$f(vC-H)_{RI}$	5.38	5.39	
$f(\nu C-H)_{R2}$	5.22	5.23	5.266
f(vC-Cl)	3.28	3.28	3.304#
$f(\delta NO_2)$	1.61	1.60	
$f(\delta O - H)$	0.72	1.04	

 Table 8. Scaled internal force constants for conformers C1 and C2 of antiviral niclosamide in gas phase by using the hybrid B3LYP level of theory.

Units are mdyn Å⁻¹ for stretching and mdyn Å rad⁻² for angle deformations, ^aThis work, ^bFrom Ref [65], [#]For 2-(4-chlorophenyl)-4H-3,1-benzoxazin-4-one

Thus, in C1, those force constants are slightly higher than the observed in C2 while a higher difference in C2 it is evidenced in the $f(\delta O-H)$ force constant, as compared with C1. This diminishing in the value can be due to that in C1; the bond C14-O3-H29 angle is not involved in the H bond formation. In contrast, in C2 with the formation of C11-O4…H23 bond that angle participle in this new H bond. When the values are compared with the predicted for 2-(4 nitrophenyl)-4H-3,1-benzoxacin-4-one) higher values in the force constants are observed for that derivative. Such differences can be easily attributed to the presence of two deactivating groups in niclosamide. In comparison, in that derivative in the nitrophenyl ring, there is only the NO₂ group different from niclosamide that presents two deactivating groups (NO₂ and Cl group).

3.8. Molecular docking studies.

According to Wu et al. [66], the niclosamide molecule has been demonstrated to have an anti-SARS-CoV activity. For that, in the present paper, we tested with COVID-19; i.e., we performed the molecular docking characteristics of NCL C_1 (in the gas phase) and C_2 (in gas and water solution) against COVID-19 protein (code: 6LU7) [67].



Figure 6. The best-docked poses of the Niclosamide C_2 (in the gas phase and water) and its conformer C1 (in the gas phase) in the COVID-19 protein.



Figure 7. H-bonding interactions between ligands and protein and their 2D interactions. (a/C₂ (in water)/COVID-19; b/ C₂ (in gas)/COVID-19 and c/ C₁ (in gas)/COVID-19.

The objective of these calculations is to provide the binding mode of these ligands with a Coronavirus containing a 3D structure. The inhibition of this protein could be an attractive target for the treatment of coronavirus. So, three complexes comprising ligands-protein were selected. In Fig. 6, we present the best-docked poses of the different ligands (having the lowest energy from the ten total poses) in the COVID-19 protein obtained by using iGEMDOCK software with the following setting: population size is 800, number 10 of generations is 80 and number of solutions is 10. The energetic results are given in Table 9, distributed into three types of non-covalent interactions: van der Waals, hydrogen-bond and electronic interactions.

Results predicted a good total energy score of the different ligands with the selected target. Docking calculations led to the following values: the total energies scores are equal to - 104.658, -97.507 and -95.362 Kcal/mol for C_2 (in water), C_2 (in the gas phase) and C_1 (in gas), respectively. It can be seen that the niclosamide C_2 in aqueous solution has the strongest binding ligand with an energy score equal to -104.658 kcal/mol. Also, it possesses the strongest

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van der Waals (-87.899 kcal/mol) and electronic interactions (0.315 kcal/mol). This indicates that the anchored inhibitors interacted strongly with C₂ in water. This finding confirms the fact that the title compound (C₂) in water is a good inhibitor against the COVID-19 better than in the gas phase. Calculations show clearly that the predominant interaction is of VDW type. For the COVID-19/C2 (in water), S-HIS-41, M-MET-49, M-HIS-164, M-MET-165, M-ARG-188, and S-GLN-189 residues formed VDW interactions; while S-HIS-41, S-TYR-54 and S-CYS-145 form H-bond interactions (as illustrated in fig. 7). In the gas phase, we notice that the total energies score of the C₂ (-71.052 kcal/mol) is greater than those calculated for C₁ (-80.049 kcal/mol). Finally and according to the docking results, we can conclude that the niclosamide molecule is a potent inhibitor against coronavirus, since it penetrates well into the active site of the COVID-19.

Target protein	Ligand	Total energy	VDW	H-bond	Electronic	Binding residue	Bond name
						S-HIS-41	H-bond
						S-TYR-54	H-bond
						S-CYS-145	H-bond
	Niclosamide					S-HIS-41	VDW
	C2 (in	-104.658	-87.899	-17.074	0.315	M-MET-49	VDW
	water)					M-HIS-164	VDW
						M-MET-	VDW
						M-ARG-	VDW
						S-GLN-189	VDW
	Niclosamide C2 (in gas)	-97.507	-71.052			M-THR-26	H-bond
						M-GLY-	H-bond
COLUD 10						M-SER-144	H-bond
COVID-19				-26.455	0	S-SER-144 M-CYS-145	H-bond
						S-CYS-145	H-bond
						S-HIS-163	H-bond
						M-LEU-141	H-bond
		-95.362				S-CYS-44	H-bond
						S-TYR-54	H-bond
						S-HIS-41	VDW
	Niclosamide C1 (in gas)		-80 049	-15 591	0.279	S-MET-49	VDW
			00.049	15.571	0.279	S-TYR-54	VDW
						M-HIS-164	VDW
						M-ARG-	VDW
						S-GLN-189	VDW

Table 9. Molecular docking energies and their distributions as non-covalent interactions.

Note that the total energy represents the sum of these three interactions. Fig. 7 (a, b and c,) shows the hydrogen bonding interactions between ligands/protein and their 2D interactions, visualized in the Discovery Studio program.

4. Conclusions

Here, the effect of medium on properties and reactivities of potential antiviral to the treatment of COVID-19, niclosamide (NCL) have been studied combining DFT calculations with molecular docking. The structures of the two most stable conformers C1 and C2 were theoretically determined in the gas phase and water, ethanol, and chloroform solvents by using

B3LYP/6-31G* calculations. The structural, electronic, topological, and vibrational properties of niclosamide were reported in the four studied media. The properties in the different solvents were studied by using the IEFPCM and universal solvation methods. The calculations in the different solutions have revealed solvation energy values for C1 and C2 in aqueous solution (ΔG_{C} = -78.43 and -64.53 kJ/mol, respectively) comparable with that observed for the antiviral agent zalcitabine (-78.92 kJ/mol). Probably, the high stability of C1, predicted by NBO studies, explains the experimental existence of C1 structure in the solid phase. Comparisons of frontier orbitals with eleven antiviral agents have evidenced the slightly high reactivity of C2 than brincidofovir, an antiviral agent used in the treatment of *Ebola* disease. Despite less negative solvation energy values of C1 and C2 in aqueous solution, as compared with eleven antiviral agents, the presence of deactivating groups (NO₂ and Cl) in the chloro-4-nitrophenyl and hydroxybenzamide rings could explain the higher reactivities observed for the two forms of NCL in different media. Here, the harmonic force fields and force constants for both forms are reported together with the assignments of 80 vibration modes expected for the two conformers.

Molecular docking results were discussed based on the type of interaction between the ligands and protein. The studies of molecular docking have shown that the niclosamide could bind to the new 2019-nCoV protein with high affinity. The three complexes showed important total energy, which makes them an important candidate to study. Finally, we hope that our contribution can help to develop a rigorous solution for this worldwide concern.

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Conflicts of Interest

The authors declare no conflict of interest.

Supporting Information Available

Tables S1-S6 and Figures S1-S6.

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Supplementary files

Table S1. Comparison of calculated geometrical parameters for the conformer C2 of niclosamide in gas and chloroform, ethanol and water solutions compared with the corresponding experimental ones in the solid phase.

Parameters	Gas	Chloroform	Ethanol	Water	Exp ^b
		Bond length	ıs (Å)		1
C11-O4	1.240	1.242	1.243	1.244	
C14-O3	1.339	1.343	1.347	1.353	1.331
N7-C9	1.396	1.394	1.393	1.393	1.395
N7-C11	1.379	1.376	1.376	1.375	1.361
C10-C11	1.483	1.484	1.484	1.483	1.540
C12-Cl1	1.761	1.762	1.761	1.760	1.730
C20-Cl2	1.757	1.764	1.765	1.764	1.737
C15-N8	1.466	1.458	1.453	1.451	1.462
N8-O5	1.230	1.233	1.235	1.236	1.220
N8-O6	1.231	1.233	1.235	1.236	1.219
RMSD	0.024	0.025	0.026	0.026	
		Bond angle	es (°)		
N7-C11-O4	121.6	121.6	121.6	121.6	
N7-C11-C10		117.1	117.0	117.1	
C10-C11-O4	121.5	121.2	121.2	121.2	
C10-C14-O3	123.4	123.2	123.1	123.0	
C19-C14-O3	117.1	117.1	117.0	117.0	111.1
C9-N7-C11	129.0	128.7	128.6	128.6	129.1
C15-N8-O5	117.5	117.9	118.3	118.3	118.1
C15-N8-O6	117.6	118.1	118.4	118.5	118.5
O6-N8-O5	124.8	123.8	123.2	123.0	123.2
N8-C15-C17	118.7	118.7	118.8	118.7	118.2
N8-C15-C18	119.7	119.6	119.6	119.6	119.3
Cl1-C12-C9	119.9	120.1	120.1	120.2	119.3
Cl1-C12-C17	118.2	118.0	117.9	117.8	118.0
Cl2-C20-C16	119.7	119.5	119.4	119.4	119.5
Cl2-C20-C21	119.6	119.4	119.3	119.3	119.3
RMSD	1.9	1.9	1.8	1.8	
		Dihedral ang	les (°)		
C9-N7-C11-O4	-3.1	-3.2	-3.5	-2.84	
C9-N7-C11-C10	177.8	177.7	177.5	178.3	
N7-C9-C12-Cl1	-0.4	-1.1	-1.0	-0.9	-3.6
O6-N8-C15-C18	179.9	-179.0	179.1	178.4	179.9
N7-Cl1- C10-C14	167.7	168.5	167.3	167.6	
C11-C10-C14-O3	2.1	1.5	1.5	1.5	
Cl2-C20-C16-C10	179.6	179.6	179.6	179.7	177.8
O4-C11-C10-C14	-11.3	-10.5	-11.5	-11.1	
RMSD	2.1	207.2	1.9	2.1	

This work, ^bRef [37].

Conformer C1								
		МК				Mulli	ken	
Atoms	gas	Chloroform	Ethanol	Water	gas	Chloroform	Ethanol	Water
1 Cl	-0.065	-0.064	-0.063	-0.062	0.000	0.000	0.000	0.000
2 Cl	-0.072	-0.075	-0.080	-0.074	-0.006	-0.009	-0.010	-0.009
3 O	-0.562	-0.577	-0.569	-0.555	-0.671	-0.671	-0.670	-0.672
4 O	-0.490	-0.489	-0.489	-0.496	-0.507	-0.509	-0.511	-0.513
5 O	-0.393	-0.391	-0.397	-0.397	-0.395	-0.396	-0.397	-0.396
6 O	-0.393	-0.390	-0.397	-0.397	-0.399	-0.400	-0.400	-0.400
7 N	-0.475	-0.485	-0.452	-0.451	-0.805	-0.806	-0.803	-0.805
8 N	0.652	0.640	0.654	0.654	0.376	0.368	0.362	0.360
		BO				ME	Р	
Atoms	gas	Chloroform	Ethanol	Water	gas	Chloroform	Ethanol	Water
1 Cl	1.219	1.218	1,218	1,219	-64.349	-64.349	-64.348	-64.348
2 Cl	1.200	1.194	1,193	1,194	-64.364	-64.364	-64.364	-64.364
30	1.931	1.936	1,937	1,936	-22.229	-22.227	-22.226	-22.226
4 O	2.004	2.002	1,998	1,996	-22.322	-22.323	-22.324	-22.324
5 O	2.041	2.037	2,034	2,033	-22.292	-22.294	-22.296	-22.296
6 O	2.036	2.032	2,029	2,028	-22.293	-22.294	-22.295	-22.296
7 N	3.251	3.253	3,254	3,255	-18.282	-18.282	-18.282	-18.282
8 N	3.974	3.973	3,973	3,972	-18.109	-18.111	-18.112	-18.112
			Сс	onformer (22			
		МК				Mulli	ken	
Atoms	gas	Chloroform	Ethanol	Water	gas	Chloroform	Ethanol	Water
1 Cl	-0.036	-0.035	-0.034	-0.034	0.016	0.016	0.017	0.017
2 Cl	-0.082	-0.090	-0.090	-0.089	-0.012	-0.016	-0.016	-0.016
3 O	-0.564	-0.564	-0.563	-0.564	-0.641	-0.643	-0.645	-0.647
4 O	-0.541	-0.541	-0.539	-0.538	-0.568	-0.570	-0.568	-0.569
5 O	-0.396	-0.395	-0.390	-0.390	-0.392	-0.392	-0.392	-0.392
6 O	-0.398	-0.396	-0.390	-0.390	-0.395	-0.395	-0.395	-0.395
7 N	-0.514	-0.515	-0.513	-0.504	-0.768	-0.765	-0.763	-0.764
8 N	0.686	0.678	0.654	0.652	0.379	0.370	0.364	0.362
		BO				ME	Р	
Atoms	gas	Chloroform	Ethanol	Water	gas	Chloroform	Ethanol	Water
1 Cl	1.227	1.225	1,226	1,226	-64.336	-64.336	-64.336	-64.336
2 Cl	1.194	1.189	1,188	1,189	-64.361	-64.362	-64.362	-64.361
3 O	1.975	1.972	1,969	1,965	-22.272	-22.273	-22.273	-22.275
4 O	1.985	1.985	1,984	1,984	-22.296	-22.296	-22.296	-22.296
5 O	2.046	2.042	2,039	2,038	-22.283	-22.285	-22.286	-22.286
6 O	2.042	2.038	2,035	2,034	-22.284	-22.285	-22.286	-22.286
7 N	3.262	3.263	3,263	3,264	-18.260	-18.258	-18.258	-18.257
8 N	3.975	3.974	3,973	3,973	-18.100	-18.101	-18.102	-18.103

Table S2. Mulliken, Merz-Kollman and NPA charges (a.u.), molecular electrostatic potentials (MEP) (a.u.) and bond orders, expressed as Wiberg indexes of two conformers of antiviral Niclosamide in different media by using B3LYP/6-31G* level of theory.

	B3LYP/6-31	G*a		
	Conformer	C1	D .1 1	XX 7 .
	gas	Chloroform	Ethanol	Water
$\pi C9 - C12 \rightarrow \pi^* C13 - C18$	59.02	58.69	58.52	58.56 102.50
$\pi C9 - C12 \rightarrow \pi^* C15 - C17$	101.95	102.91	103.50	103.50
$\pi C10 - C16 \rightarrow \pi^* O4 - C11$	90.46	92.46	93.51	94.64
$\pi C10\text{-}C16 \rightarrow \pi^*C14\text{-}C19$	105.67	105.67	105.13	105.25
$\pi C10\text{-}C16 \rightarrow \pi^*C20\text{-}C21$	73.40	74.15	74.53	74.74
$\pi C13$ -C18 $\rightarrow \pi^*C9$ -C12	101.16	100.86	100.91	100.86
$\pi C13$ -C18 $\rightarrow \pi^*C15$ -C17	79.13	78.54	78.25	78.33
$\pi C14-C19 \rightarrow \pi^*C10-C16$	66.25	65.84	65.88	65.79
$\pi C14\text{-}C19 \rightarrow \pi^*C20\text{-}C21$	99.07	99.65	100.03	99.99
$\pi C15\text{-}C17 \rightarrow \pi^*O5\text{-}N8$	117.83	121.89	125.19	126.07
$\pi C15\text{-}C17 \rightarrow \pi^*C9\text{-}C12$	71.65	72.06	72.31	72.40
$\pi C15$ - $C17 \rightarrow \pi^*C13$ - $C18$	87.19	86.61	86.23	86.15
$\pi C20$ - $C21 \rightarrow \pi^*C10$ - $C16$	101.11	102.08	102.41	102.62
$\pi C20$ - $C21 \rightarrow \pi^*C14$ - $C19$	70.73	71.06	71.48	71.65
$\Delta E_{\pi \to \pi^*}$	1224.61	1232.47	1237.87	1240.54
$\pi O5-N8 \rightarrow LP(3)O6$	50.37	50.37	50.33	50.16
$\Delta E_{\pi \rightarrow LP}$	50.37	50.37	50.33	50.16
$LP(3)Cl1 \rightarrow \pi^*C9-C12$	46.15	45.98	46.06	46.36
$LP(3)Cl2 \rightarrow \pi^*C20-C21$	49.28	47.82	47.48	47.61
$LP(2)O3 \rightarrow \pi^*C14-C19$	113.24	116.08	116.71	115.41
$LP(3)O6 \rightarrow \pi^*(1)O5-N8$	676.99	673.73	671.14	669.26
$LP(1)N7 \rightarrow \pi^*O4-C11$	252.01	252.26	254.14	255.44
$LP(1)N7 \rightarrow \pi^*C9-C12$	182.00	184.92	185.88	185.55
$\Delta E_{LP \to \pi^*}$	1319.67	1320.80	1321.42	1319.63
$LP(1)O3 \rightarrow \sigma^{*}(1)N7-H22$	48.74	52.04	50.83	53.42
$LP(2)O4 \rightarrow \sigma^{*}(1) N7-C11$	106.55	105.67	104.17	103.16
$LP(2)O4 \rightarrow \sigma^{*}(1)C10$ -C11	84.35	82.93	81.43	80.63
$LP(2)O5 \rightarrow \sigma^{*}(1)O6-N8$	80.30	79.38	78.75	78.54
$LP(2)O5 \rightarrow \sigma^{*}(1)N8-C15$	52.50	51.37	50.49	50.03
$LP(2)O6 \rightarrow \sigma^{*}(1)O5-N8$	80.09	79.38	78.88	78.58
$LP(2)O6 \rightarrow \sigma^{*}(1)N8-C15$	52.17	51.12	50.24	49.78
$\Delta E_{LP \to \sigma^*}$	504.69	501.89	494.79	494.16
$\pi^*O4\text{-}C11 \rightarrow \pi^*C10\text{-}C16$	637.37	657.60	646.40	639.29
$\pi^*O5-N8 \rightarrow \pi^*C15-C17$	73.11	76.54	79.17	79.38
π^*C9 -C12 $\rightarrow \pi^*C13$ -C18	363.79	355.43	349.32	349.32
π^*C9 -C12 $\rightarrow \pi^*C15$ -C17	715.66	724.39	730.29	730.25
$\pi^*C14\text{-}C19 \rightarrow \pi^*C10\text{-}C16$	960.44	941.13	933.19	932.56
π^*C15 -C17 $\rightarrow \pi^*C13$ -C18	805.03	756.37	721.97	722.68
$\pi^*C20-C21 \rightarrow \pi^*C10-C16$	1077.10	1136.12	1167.31	1177.26
$\Delta E_{\pi^* \to \pi^*}$	4632.49	4647.57	4627.64	4630.73
ΔE_{Total}	7731.83	7753.11	7732.04	7735.22

Table S3. Main delocalization energies (in kJ/mol) of conformer C1 of antiviral Niclosamide in different media by using B3LYP/6-31G* calculations.

^aThis work

B3LYP/6-31G*a								
	Conformer	C2						
Delocalization	gas	Chloroform	Ethanol	Water				
$\pi C9$ - $C13 \rightarrow \pi^* C12$ - $C17$	73.74	73.57	73.23	73.23				
$\pi C9$ -C13 $\rightarrow \pi^*C15$ -C18	100.40	101.03	101.57	101.53				
$\pi C10\text{-}C16 \rightarrow \pi^*O4\text{-}C11$	112.57	112.94	112.11	112.44				
$\pi C10\text{-}C16 \rightarrow \pi^*C20\text{-}C21$	66.71	67.09	67.34	67.59				
$\pi C12$ -C17 $\rightarrow \pi^*C9$ -C13	81.72	81.34	81.13	81.05				
$\pi C12$ -C17 $\rightarrow \pi^*C15$ -C18	63.41	63.24	62.99	62.99				
$\pi C15$ - $C18 \rightarrow \pi^*O5$ - $N8$	104.58	108.47	111.65	112.48				
$\pi C15$ - $C18 \rightarrow \pi^*C9$ - $C13$	74.61	75.62	76.03	76.12				
$\pi C15$ - $C18 \rightarrow \pi^*C12$ - $C17$	106.84	106.09	105.67	105.50				
$\pi C20$ - $C21 \rightarrow \pi^*C10$ - $C16$	108.39	107.64	107.01	106.88				
$\Delta E_{\pi \to \pi^*}$	892.97	897.03	898.74	899.83				
$\pi O5-N8 \rightarrow LP(3)O6$	50.83	50.87	50.83	50.70				
$\pi C20$ - $C21 \rightarrow LP(1)C19$	176.61	175.85	175.56	175.56				
$\Delta E_{\pi \rightarrow LP}$	227.43	226.72	226.39	226.26				
$\pi C10$ - $C16 \rightarrow LP(1)$ * $C14$	223.96	222.50	222.25	222.21				
$\Delta E_{\pi \rightarrow LP^*}$	223.96	222.50	222.25	222.21				
$LP(3)Cl1 \rightarrow \pi^*C12$ -C17	46.15	45.85	46.02	46.11				
$LP(3)Cl2 \rightarrow \pi^*C20$ -C21	46.36	45.14	45.02	45.14				
$LP(3)O6 \rightarrow \pi^*O5-N8$	681.88	678.58	675.86	674.15				
$LP(1)N7 \rightarrow \pi^*O4-C11$	261.12	263.67	262.13	264.59				
$LP(1)N7 \rightarrow \pi^*C9$ -C13	161.31	159.68	159.84	159.76				
LP(1)C14→π*C10-C16	270.20	270.74	271.95	273.16				
$LP(1)C19 \rightarrow \pi^*C20$ -C21	403.58	399.40	396.89	396.43				
$\Delta E_{LP \to \pi^*}$	1870.59	1863.07	1857.72	1859.35				
$LP(2)O4 \rightarrow \sigma^{*}(1)O3-H29$	84.39	90.12	85.69	87.45				
$LP(2)O4 \rightarrow \sigma^{*}(1)N7-C11$	101.70	100.15	100.36	99.69				
$LP(2)O4 \rightarrow \sigma^{*}(1)C10$ -C11	51.87	50.24	50.75	50.16				
$LP(2)O5 \rightarrow \sigma^{*}(1)O6-N8$	80.26	79.38	78.75	78.50				
$LP(2)O5 \rightarrow \sigma^{*}(1)N8-C15$	53.38	52.29	51.37	50.91				
$LP(2)O6 \rightarrow \sigma^{*}(1)O5-N8$	80.17	79.42	78.88	78.58				
$LP(2)O6 \rightarrow \sigma^{*}(1)N8-C15$	53.13	52.04	51.16	50.70				
ΔElp→σ*	504.90	503.65	496.96	496.00				
$LP(2)O3 \rightarrow LP(1)*C14$	308.94	307.73	301.59	295.78				
$\Delta E_{LP \rightarrow LP^*}$	308.94	307.73	301.59	295.78				
$\pi^*O4\text{-}C11 \rightarrow \pi^*C10\text{-}C16$	439.03	418.46	419.13	412.82				
$\pi^*O5-N8 \rightarrow \pi^*C15-C18$	79.96	83.39	85.94	86.28				
$\pi^*C12\text{-}C17 \rightarrow \pi^*C15\text{-}C18$	1003.41	1045.59	1079.44	1084.38				
$\Delta E_{\pi^* \to \pi^*}$	1522.40	1547.44	1584.51	1583.47				
ΔE_{Total}	5551.19	5568.14	5588.16	5582.9				

Table S4. Main delocalization energies (in kJ/mol) of conformer C2 of antiviral Niclosamide in different media by using B3LYP/6-31G* calculations.

^aThis work

B3LYP/6-31G* Method								
GAS PHASE								
Parameter [#]	O4-H23	RCPN1	O3-H22	RCPN2	H22-Cl1	RCPN3	RCP1	RCP2
ρ(r)	0.0219	0.0127	0.0306	0.0121	0.0159	0.0151	0.0199	0.0199
$\nabla^2 \rho(\mathbf{r})$	0.0776	0.0727	0.1119	0.0772	0.0664	0.0820	0.1564	0.1569
λ1	-0.0253	-0.0097	-0.0426	-0.0090	-0.0154	-0.0125	-0.0147	-0.0148
λ2	-0.0226	0.0265	-0.0413	0.0371	-0.0100	0.0145	0.0806	0.0797
λ3	0.1255	0.0559	0.1958	0.0491	0.0918	0.0799	0.0903	0.0919
$ \lambda 1 /\lambda 3$	0.2016	0.1735	0.2176	0.1833	0.1678	0.1564	0.1628	0.1610
Distances	2.117		1.870		2.445			
			CHI	LOFOROR	M			
Parameter [#]	O4-H23	RCPN1	O3-H22	RCPN2	H22-Cl1	RCPN3	RCP1	RCP2
ρ(r)	0.0214	0.0126	0.0319	0.0124	0.0159	0.0151	0.0199	0.0199
$\nabla^2 \rho(\mathbf{r})$	0.0757	0.0717	0.1172	0.0795	0.0663	0.0817	0.1560	0.1564
λ1	-0.0246	-0.0096	-0.0453	-0.0094	-0.0153	-0.0124	-0.0147	-0.0148
λ2	-0.0218	0.0258	-0.0438	0.0381	-0.0098	0.0142	0.0803	0.0799
λ3	0.1221	0.0554	0.2063	0.0508	0.0915	0.0798	0.0903	0.0913
$ \lambda 1 /\lambda 3$	0.2015	0.1733	0.2196	0.1850	0.1672	0.1554	0.1628	0.1621
Distances	2.130		1.851		2.448			
			E	THANOL				
Parameter [#]	O4-H23	RCPN1	O3-H22	RCPN2	H22-Cl1	RCPN3	RCP1	RCP2
ρ(r)	0.0211	0.0125	0.0322	0.0126	0.0159	0.0151	0.0199	0.0198
$\nabla^2 \rho(\mathbf{r})$	0.0747	0.0711	0.1179	0.0810	0.0663	0.0817	0.1559	0.1562
λ1	-0.0241	-0.0095	-0.0460	-0.0094	-0.0152	-0.0124	-0.0146	-0.0147
λ2	-0.0212	0.0254	-0.0441	0.0386	-0.0098	0.0142	0.0803	0.0798
λ3	0.1200	0.0552	0.2081	0.0516	0.0914	0.0799	0.0902	0.0910
$ \lambda 1 /\lambda 3$	0.2008	0.1721	0.2210	0.1822	0.1663	0.1552	0.1619	0.1615
Distances	2.139		1.849		2.449			
				WATER				
Parameter [#]	O4-H23	RCPN1	O3-H22	RCPN2	H22-Cl1	RCPN3	RCP1	RCP2
ρ(r)	0.0211	0.0126	0.0326	0.0126	0.0159	0.0151	0.0199	0.0198
$\nabla^2 \rho(\mathbf{r})$	0.0750	0.0712	0.1198	0.0810	0.0664	0.0816	0.1560	0.1562
λ1	-0,0242	-0,0095	-0,0468	-0,0095	-0,0152	-0,0124	-0,0147	-0,0147
λ2	-0,0213	0,0254	-0,0451	0,0387	-0,0097	0,0140	0,0804	0,0798
λ3	0,1205	0,0552	0,2117	0,0517	0,0913	0,0800	0,0903	0,0910
$ \lambda 1 /\lambda 3$	0,2008	0,1721	0,2211	0,1838	0,1665	0,1550	0,1628	0,1615
Distances	2.138		1.841		2.450			

Table S5. Analysis of the Bond Critical Points (BCPs) and Ring critical point (RCPs) of conformer C1 of antiviral Niclosamide in different media by using the B3LYP/6-31G* method.

*Parameters in a.u.

B3LYP/6-31G* Method										
GAS PHASE										
Parameter	04-	RCPN1	04-	RCPN2	H22-	RCPN	H22-	RCPN	RCP1	RCP2
ρ(r)	0.020	0.0121	0.047	0.0180	0.017	0.0161	0.013	0.0127	0.019	0.019
$\nabla^2 \rho(\mathbf{r})$	0.072	0.0688	0.154	0.1160	0.070	0.0910	0.060	0.0710	0.156	0.153
λ1	-	-0.0091	-	-0.0160	-	-0.0134	-	-0.0093	-	-
λ2	-	0.0249	-	0.0591	-	0.0207	-	0.0113	0.081	0.079
λ3	0.115	0.0529	0.313	0.0728	0.101	0.0835	0.083	0.0690	0.089	0.089
$ \lambda 1 /\lambda 3$	0.199	0.1720	0.254	0.2198	0.175	0.1605	0.168	0.1348	0.163	0.160
Distances	2.152		1.701		2.395		1.921			
				CHLO	OFOROR	М				
Parameter	04-	RCPN	O4-	RCPN	H22-	RCPN	H22-	RCPN	RCP1	RCP2
ρ(r)	0.0196	0.0122	0.049	0.0182	0.0169	0.0155	0.013	0.0126	0.019	0.019
$\nabla^2 \rho(\mathbf{r})$	0.0704	0.0686	0.159	0.1174	0.0674	0.0864	0.059	0.0704	0.156	0.154
λ1	-	-0.0087	-	-0.0161	-	-0.0128	-	-0.0091	-	-
λ2	-	0.0235	-	0.0603	-	0.0183	-	0.0113	0.081	0.080
λ3	0.1113	0.0538	0.326	0.0732	0.0963	0.080	0.082	0.0681	0.089	0.088
$ \lambda 1 /\lambda 3$	0.1977	0.1617	0.256	0.2199	0.1724	0.1600	0.168	0.1336	0.163	0.161
Distances	2.174		1.686		2.424		1.922			
				ET	HANOL					
Parameter	04-	RCPN	O4-	RCPN	H22-	RCPN	H22-	RCPN	RCP1	RCP2
ρ(r)	0.0192	0.0122	0.048	0.0180	0.0167	0.0154	0.013	0.0126	0.019	0.019
$\nabla^2 \rho(\mathbf{r})$	0.0692	0.0681	0.155	0.1156	0.0668	0.0854	0.059	0.0704	0.156	0.154
λ1	-	-0.0086	-	-0.0159	-	-0.0127	-	-0.0090	-	-
λ2	-	0.0228	0.079	0.0592	-	0.0176	-	0.0110	0.081	0.080
λ3	0.1087	0.0538	0.316	0.0722	0.0950	0.0804	0.081	0.0683	0.089	0.088
$ \lambda 1 /\lambda 3$	0.1969	0.1599	0.255	0.2202	0.1716	0.1580	0.166	0.1318	0.163	0.162
Distances	2.187		1.698		2.431		1.928			
				W	ATER					
Parameter	04-	RCPN	04-	RCPN	H22-	RCPN	H22-	RCPN	RCP1	RCP2
ρ(r)	0.0193	0.0122	0.048	0.0180	0.0168	0.0155	0.013	0.0126	0.019	0.019
$\nabla^2 \rho(\mathbf{r})$	0.0696	0.0682	0.157	0.1158	0.0670	0.0860	0.060	0.0709	0.156	0.154
λ1	-	-0.0087	-	-0.0159	-	-0.0127	-	-0.0092	-	-
λ2	-	0.0230	-	0.0596	-	0.0178	-	0.0116	0.081	0.080
λ3	0.1096	0.0539	0.320	0.0721	0.0955	0.0807	0.083	0.0685	0.089	0.088
$ \lambda 1 /\lambda 3$	0.1962	0.1614	0.255	0.2205	0.1728	0.1574	0.169	0.1343	0.163	0.162
Distances	2.183		1.692		2.428		1.918			

Table S6. Analysis of the Bond Critical Points (BCPs) and Ring critical point (RCPs) of conformer C2 of antiviral Niclosamide in different media by using the B3LYP/6-31G* method.

[#]Parameters in a.u.



Figure S1. Magnitudes and positions of vectors of dipole moments of conformers C1 (upper) and C2 (bottom) of niclosamide in gas phase, chloroform, ethanol and water solvents by using the $B3LYP/6-31G^*$ method.



Figure S2. Molecular structures of eleven antiviral agents compared with the corresponding to most stable conformer C2 of niclosamide.



Figure S3. Variations of atomic MK and Mulliken charges of conformers C1 and C2 of niclosamide in gas phase, chloroform, ethanol and water solvents by using the B3LYP/6-31G* method.



Figure S4. Calculated electrostatic potential surfaces on the molecular surfaces of C1 and C2 of niclosamide in gas phase. Color ranges ± 0.0463 a.u. B3LYP functional and 6-31G* basis set. Isodensity value of 0.004.



Figure S5. Molecular graphics of conformers C1 (upper) and C2 (bottom) of antiviral agent niclosamide in gas phase showing the geometry of all their bond critical points (BCPs) and ring critical points (RCPs) at the B3LYP/6-31G* level of theory.



Figure S6. Calculated gap values for C1 and C2 of niclosamide in gas phase and in chloroform, ethanol and aqueous solutions by using the B3LYP/6-31G* method.



Figure S7. Predicted electronic spectra of C1 and C2 of niclosamide in chloroform, ethanol and aqueous solutions by using the B3LYP/6-31G* method.