

Tautomerism in drugs with benzimidazole carbamate moiety: an electronic structure analysis

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Abstract Several medicinally important compounds carry benzimidazole carbamate moiety. In the scientific literature, these molecules are represented in different tautomeric forms. In this report, conformational and tautomeric preferences were analyzed on the model benzimidazole carbamate (carbendazim), so as to understand the potential energy surface of the title compounds. Quantum chemical calculations have been performed using HF, B3LYP, and MP2 methods in gas phase and solvent phase on model benzimidazole carbamate to understand the conformational and tautomeric preferences. (1) PE surface of amide–imide tautomers, (2) electron distribution, (3) AIM analysis, (4) NBO charges, (5) 1,3-H shift, etc., have been investigated for carbendazim and its conformers. The molecular electrostatic potential (MESP) surfaces of carbendazim have been studied. Further to understand the polymorphism in benzimidazole carbamate, analysis of dimers of carbendazim has been carried out. The results indicate that a neglected tautomer is important and the tautomeric equilibrium is quite subtle in these systems and it should be extensively considered in all studies related to these drugs.

Keywords Benzimidazole carbamate derivatives · Ab initio study · Tautomerism · Conformational polymorphism

1 Introduction

Benzimidazole carbamate (BC) derivatives are important class of compounds that have extensive medicinal applications in the therapy of several parasitic infections [1–3]. Thiabendazole (1), is an anthelmintic drug, introduced to treat parasitic infections. However, thiabendazole is found to have toxic effects, which obligated the researchers to develop benzimidazole carbamate derivatives. Albendazole (2), Mebendazole (3), Fenbendazole (4), Ciclobendazole (5), Flubendazole (6), Oxibendazole (7) are the relatively safe benzimidazole carbamates, which are most widely used for the treatment for a variety of parasitic infections such as hydatidosis, neurocystercosis, schistosomiasis, and intestinal nematodes (Fig. 1). All these molecules are well-known anthelmintic drugs beneficial to control, prevent, and treat against diverse set of infections caused by roundworms, hookworms, whipworms, and tapeworms [4–9]. Thus, these anthelmintic drugs are used in humans and animals for the therapy of trematode, cestode, and nematode parasitic infestations [10]. Recent reports suggested that a few of these drugs found application in cancer treatment [11]. Carbendazim (8) and Benomyl (9), which also share the benzimidazole carbamate moiety, have been shown to exhibit significant activity against fungal infections (Fig. 2) [12–15].

Low aqueous solubility (<6 mg/mL) of these molecules has restricted their formulation development in the forms of suspensions, paste, powder, or intraruminal bolus, which compels the administration in larger doses to achieve the desired therapeutic response [16–20]. Solid-state researchers working in the molecular

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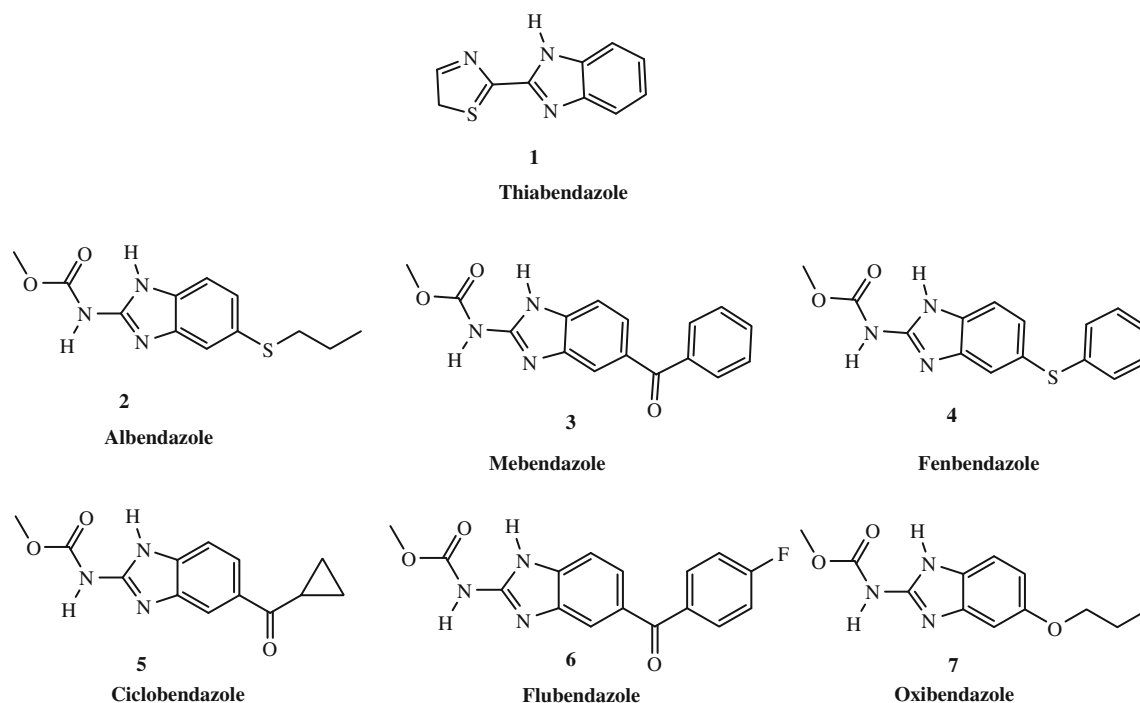


Fig. 1 Anthelmintic drugs of benzimidazole carbamates

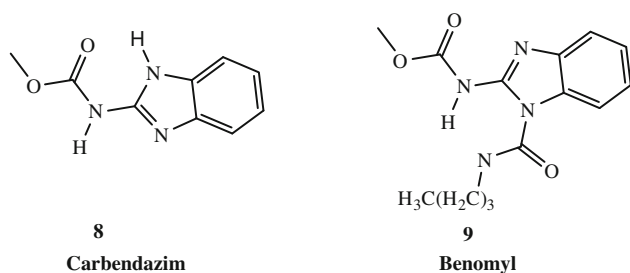


Fig. 2 Fungicidal drugs of benzimidazole carbamates

pharmaceuticals are fascinated to understand the polymorphism in the **BC** molecules, so as to improve the bio-availability profile of these molecules. In order to increase the solubility of these molecules, several strategies were adopted. Derivatives of the **BC** molecules were synthesized in the form of oxidized products and in the form of prodrugs. Recently, research is focused on the active sulfoxide metabolites of albendazole (i.e., Ricobendazole (**10**)) and fenbendazole (i.e., oxfenbendazole (**11**)) (Fig. 3) [21, 22]. Phosphonooxymethyl prodrugs of benzimidazole carbamates are reported as agents with better safety profile. Basic property of benzimidazole carbamates were exploited to prepare their acidic salts such as sulfonic acid salts. Chassaing et al. [23] reported that the intramolecular hydrogen bond present in the benzimidazole carbamate between the benzimidazole NH and the carbonyl of the carbamate moiety is the probable cause of the lack of solubility of these compounds.

In the scientific literature, benzimidazole carbamates are represented with **A** [24] and **B** tautomeric representations (Fig. 4). Crystal structure details of most **BC** drugs suggest that **B** is the correct representation [17, 20]. Sokol et al. [25] suggested that two tautomers (**B** and **C**) are possible for carbendazim (**CM**). The energy profile of tautomeric process in this class of drugs was not explored. Pranzo et al. recently reported the crystal structure of albendazole, in which a polymorph with the tautomeric representation **C** is found to be more stable. Electronically, tautomer **A** suffers from lone pair-lone pair repulsion between imidazole nitrogen and carboxylic oxygen, whereas conformation **B** represents a tautomer, which is devoid of such repulsion, and in addition, it carries stabilizing intramolecular hydrogen bonding interaction. Structure **C** represents the imide form of benzimidazole (*N*-heterocyclic carbene) moiety; this tautomer also maintains the stable hydrogen bonding interaction. Thus, the critical phenomenon of structural tautomerism observed in the **BC** molecules prompted us to explore the PE surface of benzimidazole carbamate (**BC**) derivatives quantum chemically, to identify the conformational and tautomeric preferences of these molecules.

Understanding the atomic level information regarding the drug molecules helps the pharmaceutical scientist in the design of new molecules and development (so as to overcome the shortcomings associated with drugs) of the existing drug molecules in different areas of pharmaceutical research—(1) Molecular pharmaceuticals scientists to generate new polymorphs to improve the solubility and

Fig. 3 Metabolites of albendazole (ricobendazole) and fenbendazole (oxefenbendazole)

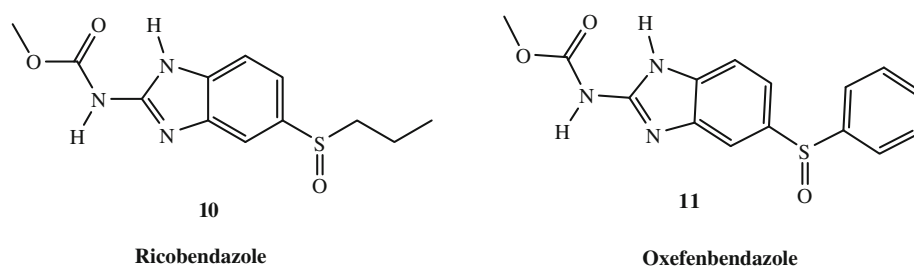
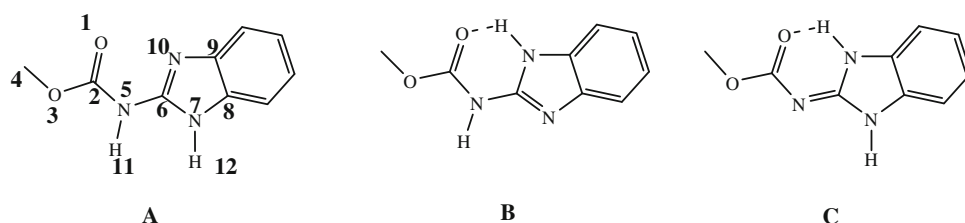


Fig. 4 Three important tautomers of benzimidazole carbamate (carbendazim)



bioavailability profile of drugs, (2) Computational chemists to understand the drug receptor interactions that help in the design of new potent molecules, (3) Medicinal chemists and pharmacologists to identify the stereo and electronic features that are responsible for drug resistance.

Several groups highlighted the role of tautomeric preferences of drugs [26–32]. Our group reported the detailed electronic structure studies, conformational and tautomeric preferences on aminoguanidine, biguanide, guanyltiourea, sulfonylurea, and guanylurea derivatives, which are therapeutically important [33–41]. Lipkowitz and McCracken performed semi-empirical (MNDO) studies on albendazole, oxbendazole, etc., to understand the conformational tautomerism in these molecules [42–45]. They suggested that the conformer **B** is the most stable molecular model, but did not consider structure **C** in the study.

In the current report, an electronic structure analysis and tautomerism details of the model benzimidazole carbamate, Carbendazim (**CM**) are presented. To understand the influence of solvent medium on the conformational preferences, implicit solvent models were employed. To comprehend the polymorphism in **BC** molecules, analysis on the dimers has been carried out. The results indicate that the tautomerism is highly probable in this class of compounds, a tautomer which was generally ignored in the chemistry literature turns out to be quite important.

2 Methodology

Ab initio molecular orbital (MO) [46, 47] and density functional theory (DFT) [48, 49] calculations have been carried out using the GAUSSIAN03 software package [50]. Complete optimizations have been performed on various tautomers and rotamers of carbendazim (**CM-1** to **CM-9**, Fig. 5), transition states, and various dimeric arrangements of

conformers of the model **BC** moiety, using HF (Hartree–Fock), B3LYP (Becke3, Lee, Yang, Parr), [51, 52] and MP2 (Moeller-Plesset perturbation) [53, 54] methods with the 6-31+G(d) basis set. 6-31+G(d) basis set incorporates the polarization function and diffuse functions, and hence it is sufficiently rigorous in tautomerism studies as evident from the published work from our group [37–39, 41]. Solvent level optimization studies have been performed using the integral equation formalism versions of the polarizable continuum model (IEFPCM; using $R_{\text{MIN}} = 0.5$, $\text{OFAC} = 0.8$) method [55] at 6-31+G(d) basis set to understand the influence of the solvent medium. A few representative drugs of benzimidazole carbamate derivatives have also been optimized using the B3LYP/6-31+G(d) method to understand the differences in the conformational preferences of these drugs. Intramolecular hydrogen bonding is confirmed by AIM (atoms in molecules) calculations using AIM2000 software package [56, 57]; NBO (natural bond orbital) analysis has been employed in the estimation of partial atomic charges [58, 59]. NICS (nuclear independent chemical shift) [60, 61] values were obtained using the GIAO method [62] using B3LYP/6-31+G(d) geometries to get the extent of electronic delocalization in the ring-forming conformations. The energy and geometric parameters used in the discussion are from the B3LYP/6-31+G(d) method unless otherwise specifically mentioned. MESP calculations have been carried out to know the surface properties of these compounds [63].

3 Result and discussion

3.1 Electronic structure and conformational analysis

Carbendazim (**CM**) can be taken as a representative of benzimidazole carbamate (**BC**)-based drugs. Considering the chemically feasible 3D geometries, 15 conformers and

tautomers were generated for **CM**. Semi-empirical PM3 method was initially used to optimize the generated models, high energy conformers were excluded, and nine important tautomers/conformers were selected for further

accurate quantum chemical (HF, B3LYP, and MP2 methods) analysis. These studies lead to the recognition of nine important tautomers/rotamers (**CM-1** to **CM-9**) of carbendazim as specified in Table 1. Figure 5 shows the 3D

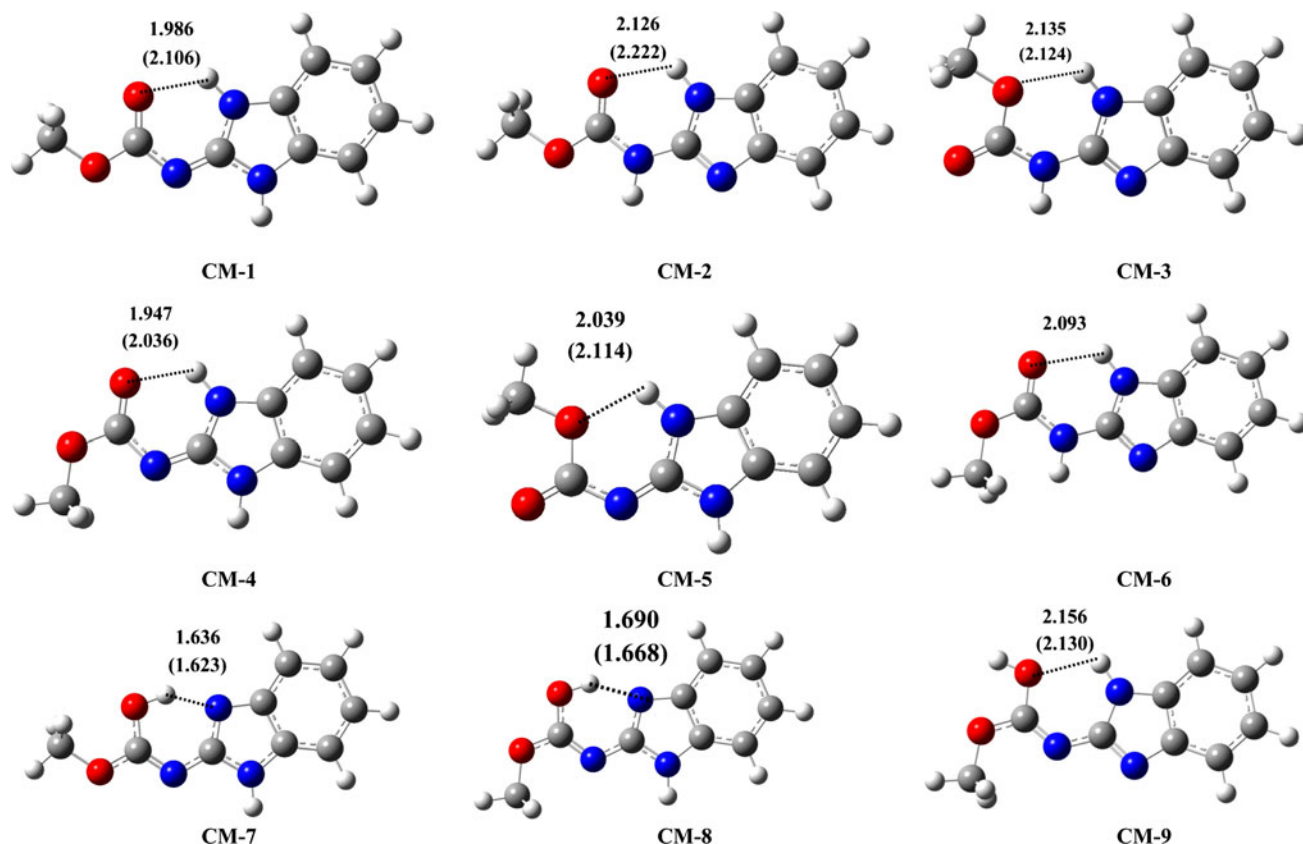


Fig. 5 Optimized 3D geometries of benzimidazole carbamate (carbendazim) **CM-1** to **CM-9**. The H-bond lengths obtained using B3LYP/6-31+G(d) method are given, data in parentheses corresponding to water medium

Table 1 Relative energies (in kcal/mol) of various tautomers of model benzimidazole carbamate moiety carbendazim (**CM**)

Conformer	Structural description	Gas phase			Solvent phase ^a	
		HF	MP2	B3LYP	Water B3LYP	Methanol B3LYP
CM-1	N5-N7 H shift (1,3-H shift tautomer of CM-2)	0.00	0.00	0.00	0.00	0.00
CM-2	Generally referred CM structure	−3.31	−5.46	−2.47	0.05	0.08
CM-3	C2-N5 rotamer of CM-2	0.51	−2.29	1.01	4.48	2.35
CM-4	O3-C4 rotamer of CM-1	2.80	2.12	2.27	3.38	1.35
CM-5	C2-N5 rotamer of CM-1	6.30	4.82	5.04	2.07	2.53
CM-6	O3-C4 rotamer of CM-2	6.18	2.89	5.27	–	6.98
CM-7	N10-N1 H shift (1,5-H shift tautomer of CM-1)	13.90	8.22	9.70	15.47	13.18
CM-8	1,5-H shift of CM-4	11.73	6.51	8.70	15.00	12.89
CM-9	1,3-H shift of CM-4	22.97	17.26	20.16	23.99	21.79
CM1-TS	Transition state for 1,3-H shift in CM-2	55.91	41.46	42.45	51.75	49.42

Basis set used for all optimizations is 6-31+G(d)

All relative energies are corrected for zero-point vibrational energy and thermal correction to Gibbs free energy

The ZPE and Gibbs free energy corrected absolute energy values are given as supporting information in Table S1 and S2

^a Implicit solvent analysis using IEFPCM level using B3LYP/6-31+G(d) method

Table 2 Tautomeric preferences observed for the **BC** drugs in both gas phase and water phase

Drugs	ΔE_g (kcal/mol)	ΔE_w (kcal/mol)
Albendazole	2.01	−0.77
Mebendazole	3.07	0.45
Fenbendazole	2.42	−0.40
Ciclobendazole	3.06	−0.16
Flubendazole	3.32	0.41
Oxibendazole	1.87	−0.45
Ricobendazole	1.03	−2.09
Oxifenbendazole	3.76	2.31

Negative value of ΔE indicates the conformational preference toward **CM-1** type tautomer, and positive value indicates the preference toward **CM-2** type tautomer

Basis set used for all optimizations is 6-31+G(d)

All relative energies are corrected for zero-point vibrational energy and thermal correction to Gibbs free energy

The ZPE and Gibbs free energy corrected absolute energy values are given as supporting information in Table S1 and S2

Implicit solvent analysis using IEFPCM level using B3LYP/6-31+G(d) method

structures of **CM-1** to **CM-9** and Table 1 shows their relative energies both in gas phase and solvation phase.

The relative energy data given in Table 1 show that there is a tautomeric equilibrium between tautomers **CM-1** and **CM-2**. Under gas-phase conditions, **CM-2** is favored, whereas in solvent conditions, **CM-1** is favored. **CM-1** suffers from lone pair (O_3)-lone pair (N_5) repulsion. The influence of this electrostatic repulsion gets reduced in polar solvents, and hence in solvent media, **CM-1** turned out to be marginally more stable. Because the ΔE between these two tautomers is quite low in water medium (0.05 kcal/mol), it can be concluded that all the drugs based on **BC** moiety may exist in tautomeric equilibrium in physiological conditions. This is further supported by the data given in Table 2.

The ΔE_g and ΔE_w values for Albendazole obtained using B3LYP/6-31+G(d) method, respectively, are 2.01 and −0.77 kcal/mol. Similarly, the comparative ΔE values of Mebendazole (ΔE_g , ΔE_w : 3.07, 0.45 kcal/mol), Fenbendazole (ΔE_g , ΔE_w : 2.42, −0.40 kcal/mol), Ciclobendazole (ΔE_g , ΔE_w : 3.06, 0.16 kcal/mol), Flubendazole (ΔE_g , ΔE_w : 3.32, 0.41 kcal/mol), Oxibendazole (ΔE_g , ΔE_w : 1.87, −0.45 kcal/mol), Ricobendazole (ΔE_g , ΔE_w : 1.03, −2.09 kcal/mol), and Oxifenbendazole (ΔE_g , ΔE_w : 3.76, 2.31 kcal/mol) clearly establish the existence of tautomeric equilibrium in these drugs. Recently, Pranzo et al. [18] reported about the **CM-1** type of polymorph experimentally; however, PE surface of the benzimidazole carbamates and the electronic structure distribution of these compounds were not comprehensively reported with

detailed quantum chemical analysis till now. Considering the importance of the tautomerism in drug molecules in the recent past [26–41], the current results on the tautomerism of benzimidazole carbamates drugs may gain importance.

Both **CM-1** and **CM-2** are characterized by intramolecular hydrogen bonds. Indeed, AIM calculations also confirmed the presence of intramolecular hydrogen bond in these species. For example in **CM-1**, a bond critical point was observed between O1 and H11–N10 indicating the presence of a hydrogen bond, which is characterized by charge density $\rho = 0.02704$, density laplacian $\nabla^2\rho = 0.09449$, and ellipticity $\varepsilon = 0.1003E + 00$. A ring critical point was also observed corresponding to the six-membered ring that is formed due to intramolecular hydrogen bond and is characterized by density 0.1656E − 01. The energy difference between **CM-2** and **CM-1** is only about 2.47 kcal/mol in gas-phase conditions. The estimated barrier for the unimolecular 1,3-H shift is quite high (44.92 kcal/mol) in **CM-1**, and hence, it is not a favorable process under regular room temperature conditions. However, the estimated barrier for the unimolecular 1,3-H shift via water-assisted mechanism (explicit one water molecule) is found to be 12.92 kcal/mol suggesting that 1,3-H shift through water is a favorable process for the inter-conversion of **CM-1** to **CM-2** and vice versa.

To elucidate the origin of the tautomeric preferences in this class of drugs, calculations have been carried out on the base moiety carbamate guanidine (**CG**, Fig. 6, Table 3). **CG-1** has been found to be most stable in gas phase (by 6.19 kcal/mol) as well as in solvent phase (by 8.57 kcal/mol) conditions. This clearly points out that on a simple guanidine, the carbamate substituent prefers to be on the imine nitrogen than the amine nitrogen. The marginal preference for **CM-2** over **CM-1** (in gas phase in comparison with the strong preference for **CG-1** over **CG-2** indicates that the aromatic ring in the imidazole ring of carbendazim is playing a role in shifting the tautomeric preferences in carbendazim, the **CM-2** carries aromatic stabilization, whereas in **CG**, this component is absent. To further verify this aspect, tautomerization in carbamate imidazoles (**CI**, Fig. 6) has been studied. The energy difference between **CI-1** and **CI-2** is 4.86 kcal/mol favoring **CI-2**. However, the energy difference is reduced to 0.42 kcal/mol favoring **CI-2** under solvent conditions. Table 3 shows tautomerization energies associated with **CG** and **CI**. NICS studies were performed to understand the aromaticity of benzimidazole, imidazole, and the delocalization of guanidine derivatives. After tautomerism (in **CM-1**), the aromatic character in five-membered ring (on *N*-heterocyclic carbene unit) is reduced. The NICS value for the five-membered ring in **CM-1** (5.78) is relatively smaller than that in **CM-2** (8.03) (See supporting information, Table S10). These data support that in

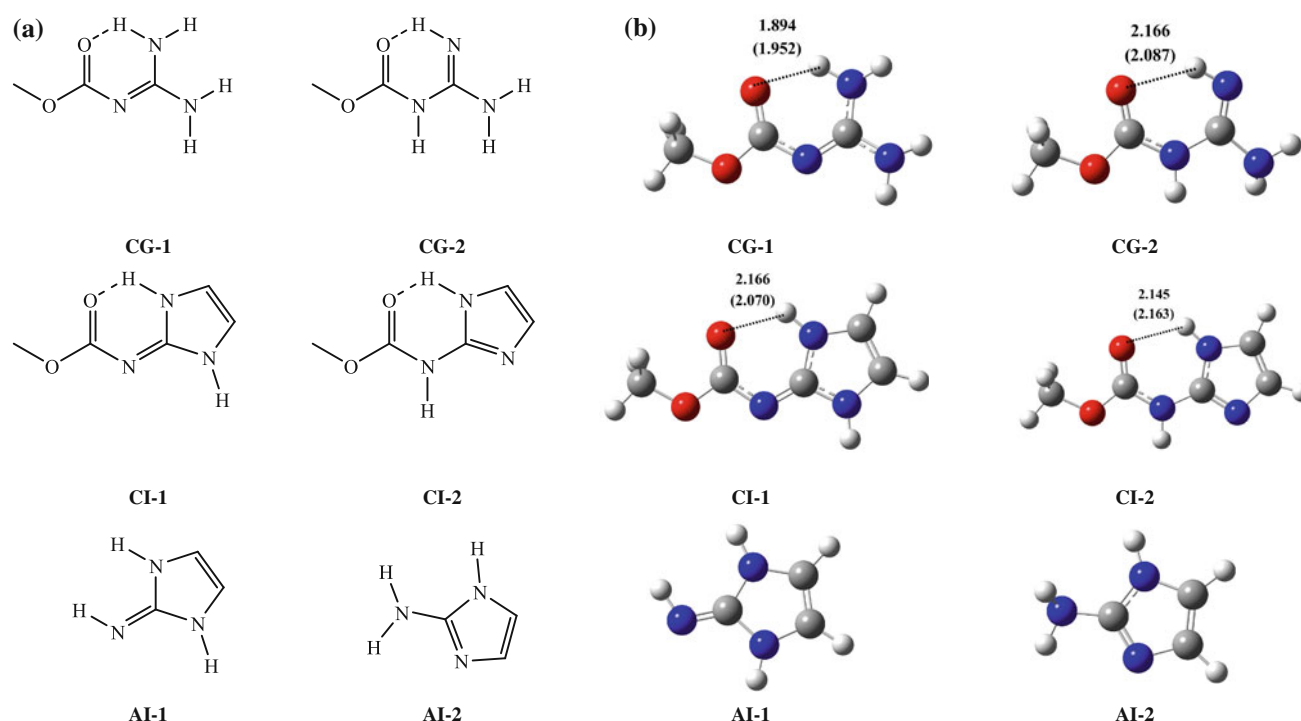


Fig. 6 **a** 2D Structures **b** Optimized 3D geometries of carbamate guanidine (CG), carbamate imidazole (CI), and amino imidazole (AI), intramolecular hydrogen bond distances are given in Å units.

Table 3 Tautomeric preferences observed for the Carbamate guanidine (CG) and Carbamate imidazole (CI) isomers, in both gas phase and water phase

Molecule	ΔE_g (kcal/mol)	ΔE_w (kcal/mol)	NBO charge on central nitrogen in CM-1 type isomer
Carbamate guanidine	−6.19	−8.57	−0.661
Carbamate imidazole	4.86	0.42	−0.658
Carbendazim	2.47	−0.05	−0.644

Negative value of ΔE indicates the conformational preference toward **CM-1** type, and positive value indicates the conformational preference toward **CM-2** type isomer

Basis set used for all optimizations is 6-31+G(d)

All relative energies are corrected for zero-point vibrational energy and thermal correction to Gibbs free energy

The ZPE and Gibbs free energy corrected absolute energy values are given as supporting information in Table S1 and S2

Implicit solvent analysis using IEFPCM level using B3LYP/6-31+G(d) method

carbendazim, the tautomerization tilts the balance toward **CM-1** but the aromatic character of the five-membered ring tilts the balance toward **CM-2**; overall, **CM-2** is marginally more stable.

On the PE surface of **CM**, one more low energy structure could be identified. This is also characterized by

The H-bond lengths obtained using B3LYP/6-31+G(d) method are given, data in parentheses corresponding to water medium

intramolecular hydrogen bond. **CM-3** is an isomer of **CM-2** in which the OMe group of carbamate is involved in intramolecular hydrogen bond. This structure may also be possible under equilibrium conditions because this tautomer is only about 1.01 kcal/mol unstable. Similarly, **CM-4** is a conformer of **CM-1**, which is only about 2.27 kcal/mol less stable than **CM-1**. These data clearly indicate that the carbendazim may adopt any of the low-lying energy states **CM-1** to **CM-4** under equilibrium conditions. **CM-5** and **CM-6** are other alternative structures whose energies lie within 6 kcal/mol with reference to **CM-1**. **CM-7** is an enol tautomer stabilized through strong intramolecular hydrogen bond. This may be obtained by 1,5-H shift in **CM-1**. This tautomer is 10 kcal/mol less stable than **CM-1**. But this particular tautomer may not be observable owing to the fact that there are alternatively many other stable possibilities, i.e., **CM-1** to **CM-4**.

Figure 7 shows the molecular electrostatic potential (MESP) of **CM-1** and **CM-2**. The electrostatic potential of **CM-1** shows the concentration of electron density at N5. Clearly, there is a large difference in the MESP of **CM-2** with respect to that of **CM-1**. In **CM-1**, negative–negative–positive electrostatic potentials due to the O–N–NH unit are noticeable, whereas in **CM-2**, alternate negative–positive–negative electrostatic potentials due to the O–NH–N unit are evident. This factor also must be contributing to the marginally greater stability of **CM-1**.

Fig. 7 Molecular electrostatic potential (MESP) surfaces of carbendazim **CM-1** and **CM-2** plotted onto a surface of constant electron density (0.002 e/au³)

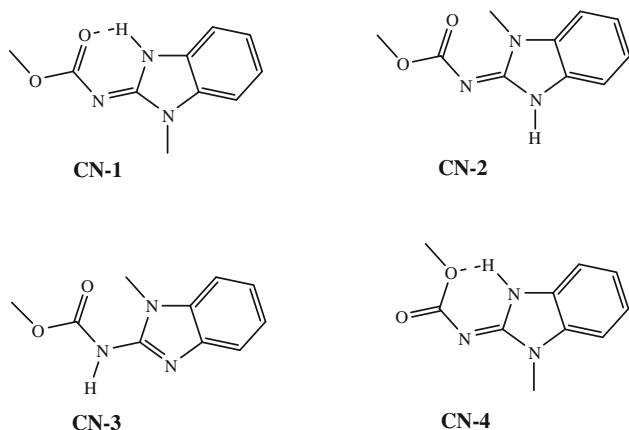
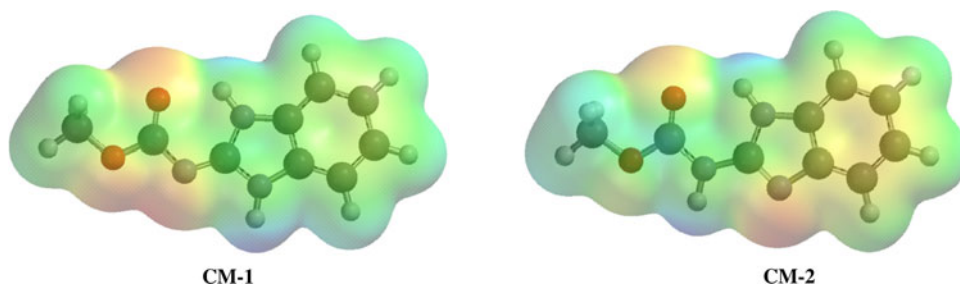


Fig. 8 2D Structures of tautomers of NSC135301

Machatha et al. [64] reported the X-ray crystal structure of carbendazim hydrochloride dehydrate. The calculated protonation energy of **CM-1** is 221.42 kcal/

mol. The protonation energy of **CM-1** is quite comparable to that of simple guanidine (235.0 kcal/mol), **CG-1** (220.58 kcal/mol) and **CI** (217.27 kcal/mol). It may be presumed that the tautomerism between **CM-1** and **CM-2** may take place through this protonated species because unimolecular tautomerism can be ruled out in the case of **BM** drugs. Deprotonation is also a possible reaction on benzimidazole ring of carbendazim. This is evidenced by the fact that albendazole is described as an amphoteric species [65]. The Gibbs free energies due to deprotonation of H11 and H12 in **CM-1**, respectively, are 327.75 and 321.99 kcal/mol at the B3LYP/6-31+G(d) level, which is slightly smaller than the deprotonation energy of acetic acid (332.66 kcal/mol). The above-reported data on protonation and deprotonation are in tune with the amphoteric character of this class of compounds.

To understand the conformational and tautomeric preferences in *N*-substituted benzimidazole derivatives, quantum chemical studies were performed using B3LYP/6-

Table 4 Relative energies (gas phase) of dimeric **CMs** (**D1–D8**) and their stabilization energies obtained using B3LYP/6-31+G (d) method

Dimer	Relative energy (kcal/mol) ^a			Molecular unit	Stabilization energy (kcal/mol) ^b
	Gas phase	Solvent phase ^c			
		Water	Methanol		
D1	0.00	0.00	0.00	CM-1	7.02
D2	−0.14	0.80	0.64	CM-2	2.22
D3	6.87	4.45	4.71	CM-5	10.22
D4	7.06	5.19	5.31	CM-3	1.97
D5	7.63	5.47	6.48	CM-2	−5.56
D6	11.01	–	8.06	CM-4	0.55
D7	11.99	–	42.67	CM-3	−2.96
D8	21.80	15.35	15.56	CM-6	−4.24
D9	23.76	27.30	27.01	CM-7	2.66
HD1	2.01	–	–	CM-7	−2.44

The ZPE and Gibbs free energy corrected absolute energy values are given as supporting information in Table S3

All relative energies are corrected for zero-point vibrational energy and thermal correction to Gibbs free energy

^a Relative energies of **D1–D9**

^b Gas-phase stabilization energy due to dimer formation from various **CMs**

^c Implicit solvent analysis using IEFPCM level using B3LYP/6-31+G(d) method

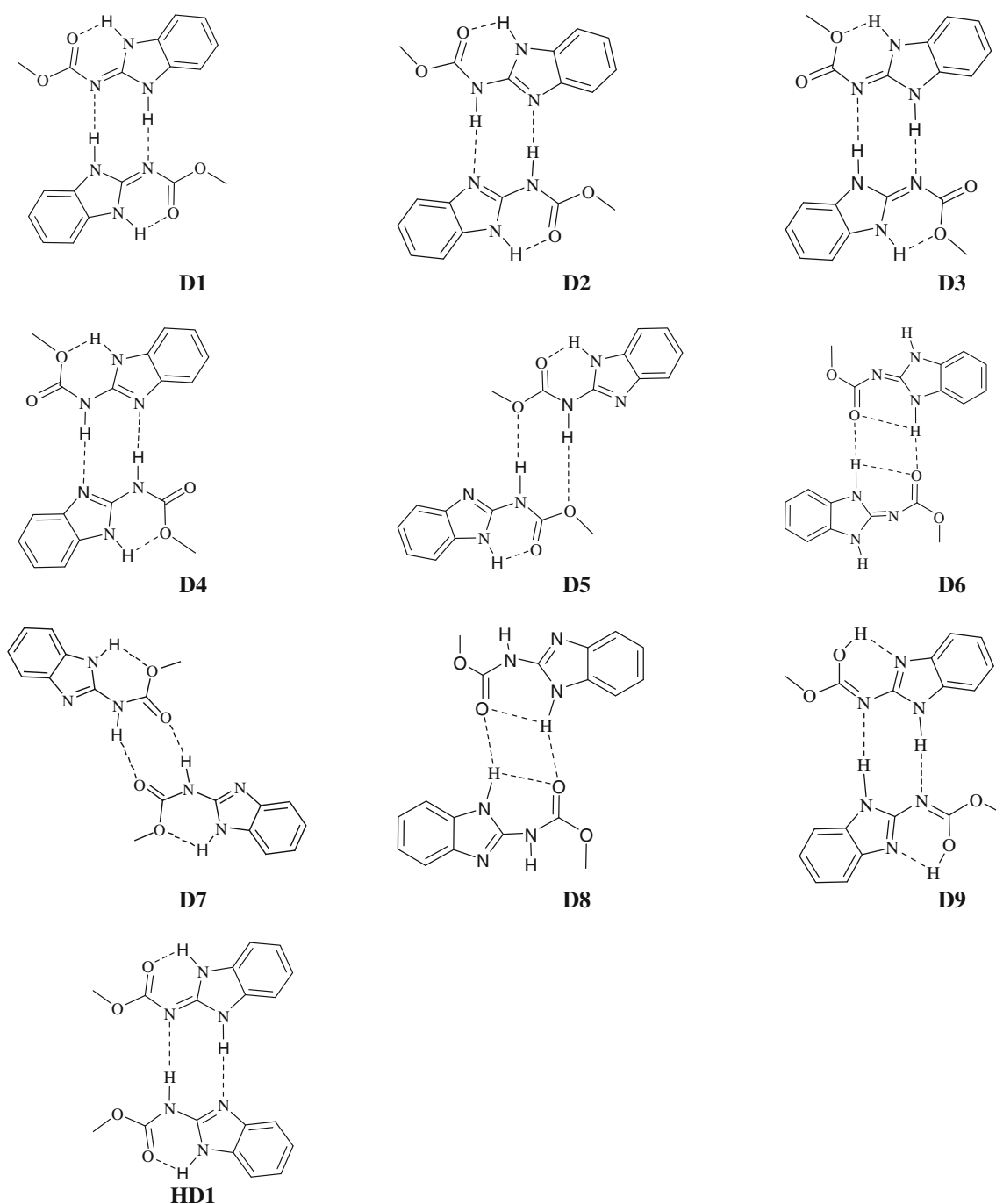


Fig. 9 Various dimeric arrangements of **D1** to **D9** with possible intermolecular interactions

31+G(d) method on NSC135301 (Fig. 8) [66]. **CN-1** was found to be more stable among possible tautomers of NSC135301, i.e., **CN-1** to **CN-4**. Both **CN-1** and **CN-4** are characterized by intramolecular hydrogen bonds. Optimized 3D geometries and the relative energy data of NSC135301 are given in supporting information (Figure S1 and Table S11).

3.2 Intermolecular interactions in carbendazim dimers and their possible contribution to polymorphism

Polymorph is ‘a solid crystalline phase of a given compound resulting from the possibility of at least two different arrangements of the molecules of that compound in the solid state’ [67]. Sokol et al. [25] reported the structure of

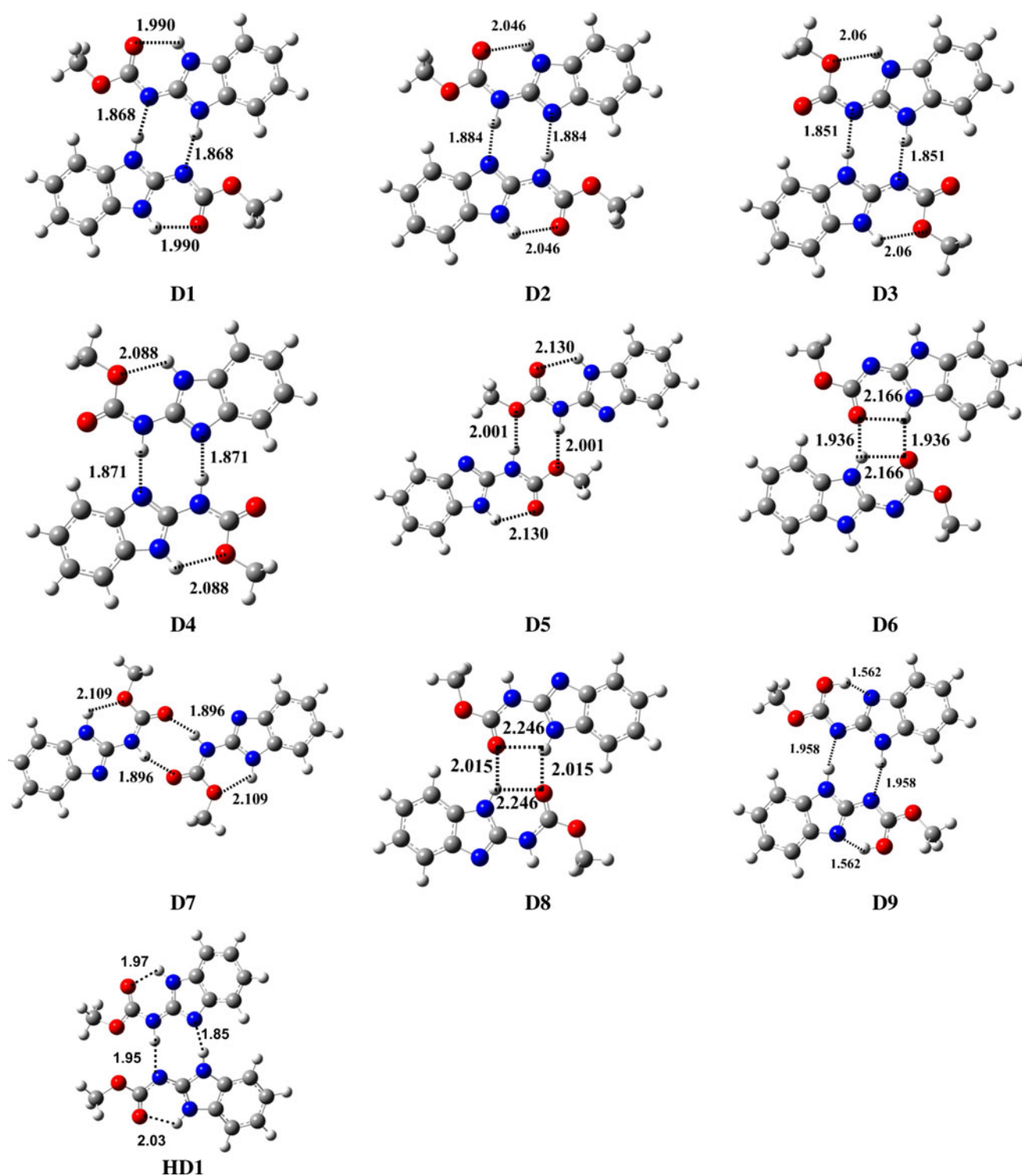


Fig. 10 Optimized 3D structures of **D1** to **D9** with inter and intramolecular hydrogen bond distances given in Å

dimer of ethyl-2-benzimidazole carbamate, in which anti-parallel arrangement of the monomeric units has shown a dimeric state with two intermolecular hydrogen bonds. Pranzo et al. [18] reported that albendazole polymorph form I is more soluble (commercial) than form II

(recrystallized in DMF; dimethyl formamide). They also reported that a polymorph of albendazole with **C** tautomer is relatively more preferable, over a polymorph with **B** tautomer for albendazole. Mebendazole is known to exist in three polymorphic forms (A, B, and C), with polymorph

C being most efficacious [19, 68, 69]. Polymorph B is reported to be toxic. Martins et al. [17] reported that the intermolecular contacts influence the conformational and geometric features of polymorph C of mebendazole. Ferreira et al. [20] reported that mebendazole (form A) can exist as a network of dimers in solid state and suggested that the dimeric state is similar to a dimer of **CM-2**. Similar dimeric structures are also possible for the tautomers **CM-2** to **CM-7**. The polymorphic states of **BM** drugs are expected to emerge from the various tautomers of these drugs and their dimeric states. To estimate the energy factors associated with the dimers of **CM-1** to **CM-7**, quantum chemical analysis using B3LYP/6-31+G(d) level has been carried out. Table 4 lists the energies of stabilization due to dimer formation (**D1-D9**) and the relative energies of the dimers, the 2D and 3D structures of the dimers **D1-D9** are shown in Figs. 9 and 10, respectively. **D1** is the most stable dimer and **D2** is next best with a relative energy of 0.80 kcal/mol in aqueous phase. **D1** is a dimer from the monomeric unit **CM-1** and **D-2** is a dimer from the monomeric unit **CM-2**. Both of them are characterized by two intermolecular hydrogen bonds. The presence of inter and intramolecular hydrogen bonds in **D1** is confirmed by performing AIM analysis. The charge density $\rho = 0.0363$, density laplacian $\nabla^2\rho = 0.0986$, and the ellipticity $\varepsilon = 0.0609$ at the intermolecular BCP in **D1** are clearly indicative of a typical hydrogen bond.

Stabilization energy due to the formation of **D1** is 7.02 kcal/mol, and the stabilization energy due to the formation of **D2** is 2.2 kcal/mol. This is the only parameter which differentiates **D1** and **D2**. On this count, it may be considered that formation of **D1** is thermodynamically preferred over the formation of **D2**. Interconversion between **D1** and **D2** may lead to two different polymorphic states of **BM** drugs. However, owing to minor changes between **D1** and **D2** and to the greater similarities in terms of geometric, thermodynamic parameters, as well as intermolecular interactions, it may not be easy to determine the polymorphic states arising from **D1** and **D2**. Since the stabilization energy in **D1** formation is relatively more favorable, it may be safely concluded that the tautomer **CM-1** is the preferred tautomer of **CM** and other **BM** drugs in solid state.

D3 is a dimer from the monomeric unit **CM-5**, and **D4** is a dimer from the monomeric unit **CM-3**. They are relatively less stable but the formation of these dimers also is thermodynamically quite feasible because of the strong stabilization energies. These dimeric structures can lead to clearly identifiable polymorphs. The dimers **D5-D9** are less stable on a relative energy scale, and these structures cannot be expected in solid-state conditions. Heterodimer (**HD-1**) was constructed from the monomers **CM-1** and **CM-2** to check the possibility of existence of heterodimer.

The stabilization energy of the complex is suggesting that the heterodimer complex is 2.44 kcal/mol (Table 4) unstable with respect to the addition of individual monomeric energies of both **CM-1** and **CM-2**.

4 Conclusions

Quantum chemical exploration of the carbendazim was performed, so as to understand tautomeric preferences of benzimidazole carbamate derivatives. Quantum chemical analysis revealed that the tautomer represented by **CM-1** structure is important. This representation was not considered in previous studies. In gas phase, **CM-1** structure is marginally less stable, whereas in solvent conditions, **CM-1** is marginally favoured and in solid-state conditions also **CM-1** structure is marginally preferred. Thus, tautomeric preferences of the benzimidazole carbamate drugs are quite environment-dependent. The preference for the **CM-1** tautomer originates from the guanidinic structure; the preference toward **CM-2** tautomer is associated with aromatic character of cyclic guanidine units. Stabilization energy due to dimer formation is 7.02 kcal/mol from **CM-1** and 2.22 kcal/mol from **CM-2**. This understanding can help in the design of new polymorphic forms, which will provide desired solubility and bioavailability properties.

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