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NMR and NBO calculation of benzimidazoles and pyrimidines: Nano physical parameters investigation

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Six theoretical methods were used for the calculation of physical parameters for derivatives of pyrimidines and benzimidazols. We used Gaussian 98 at NMR and NBO calculations by using HF method with 6-31G, 6-31G* and 6-31+G basis set and B3LYP, BLYP and B3PW91 methods with 6-31G basis set. Chemical shift was drawn curve for all of the atoms in each molecule. The thermo chemical parameters including thermal energy(ΔE), atomic charges, chemical shift anisotropy(δ), asymmetry parameter(η), chemical shift anisotropy($\Delta \sigma$), dipole orientation, isotropic, anisotropic, NMR determinant and distance matrix compounds. Also the natural bond orbital (NBO) analysis has been performed which seem informative to show some important atomic and structural features.

Key words: Benzimidazoles, pyrimidines, NBO analysis, Nano physical parameters.

INTRODUCTION

Recently, medicinal chemists have taken great interest in the combinatorial methods for the design and synthesis of pharmacologically-relevant heterocyclic molecules (Dolle and Nelson, 1999; Dolle, 2000, 2004). Inexpensive and readily-available domestic microwave ovens transform electromagnetic energy into heat, thus the absorption and transmission of the energy varies greatly from that of conventional heating. The benzimidazole nucleus is an important structure in numerous natural and synthetic compounds and in medicinal chemistry. Solidphase approaches for benzimidazole synthesis have been published (Smith et al., 1999), however the cyclization of benzimidazole catalyzed by aluminamethanesulfonic acid (AMA) under microwave irradiation has not been reported. The appropriate carboxylic acids were reacted with 1, 2 phenylenediamine in the presence of AMA under microwave irradiation to give the corresponding benzimidazoles in good to excellent yields (Scheme 1).

Structures containing benzimidazole, well-known to have a wide range of biological properties, have commercial applications in various realms of therapy, including antiulcerative, anti-hypertensive, antiviral, antifungal, antitumor and antihistaminic agents, and antihelminthic agents in veterinary medicine (Spasov et al., 1999). Benzimidazole derivatives have found the appreciation in diverse therapeutic areas including antimicrobial activity (Kus et al., 2004; Wu et al., 2003; Mollaamin et al., 2008; Ozden et al., 2005; Sztanke et al., 2006), the activity against several viruses such as HIV (Porcari et al., 1998; Mollaamin et al., 2010; Samia et al., 2006), antiallergic (Kilcig and Altanlar, 2006; Nakano et al., 1999), antioxidant, antihistaminic (Vijayakumar and Jafar Ahamed, 2010), antitubercular (Yadav and Srivastava, 2011; Kuchkguzel et al., 2001), antiasthmatic (Souness et al., 2000), antidiabetic (Senten et al., 2003; Black et al., 2005), anticancer (Pal et al., 2011; Sun et al., 2011; Kruse et al., 1989; Islam et al., 1991; Ramla et al., 2007), antitumor (Denny et al., 1990 ; Tatsuta et al., 2005), antiulcer (Jung et al., 1993; Hazelton et al., 1995), antihelmentic (Karen, 2006), HIV-1 reverse transcriptase inhibitors (Gardiner et al., 2003), anticoagulant (Young et al., 2006), anti inflammatory(Fox et al., 2009), antibacterial (Rosowsky et al., 1997; Andrzejewska et al., 2004), the series of biologically active benzimidazoles (AJafar et al.,2009). Furthermore, these heterocycles are considered to be privileged structures by medicinal chemists.

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Scheme 1. Carboxylic acids reaction with 1, 2 phenylenediamine in the presence of AMA under microwave irradiation to give the corresponding benzimidazoles in good to excellent yields.



Scheme 2. Preparation of dihydropyrimidinones through Biginelli reaction under mild conditions.



Scheme 3. Ethyl β - aminocrotonate undergoes an extremely facile reaction with phenylisocyanate or methyl isocyanate to form an intermediate ureido derivative which undergoes cyclization.

This seems to be because pyrimidines represents one of the most active class of compounds possessing wide spectrum of biological activity, namely, significant *in vitro* activity against unrelated DNA and RNA, viruses including polio herpes viruses, diuretic, antitumor, anti HIV, cardiovascular (Kappe,1993). Biginelli compounds show a diverse range of biological activities. The interest focused on Biginelli compounds leading to the development of Nitractin (A) (Marsault and Peterson, 2011), that has excellent activity against the virus of trachoma group, the same compounds also exhibit antibacterial activity. Application of dodecylphosphonic acid (DPA) (Ghassamipour and Sardarian, 2009), solid Bronsted acid, is reported for the preparation of dihydro-pyrimidinones through Biginelli reaction under mild conditions (Scheme 2).

Ethyl β- aminocrotonate undergoes an extremely facile

reaction with phenylisocyanate or methyl isocyanate to form an intermediate ureido derivative which undergo cyclization as treatment with base (Ghassamipour and Sardarian, 2009; Gupta et al., 2010) (Scheme 3).

METHODS OF COMPUTATION

In Stage 1, we started a ChemDraw and constructed molecules, after which the results were saved as a ChemDraw file. In Stage 2, this file was reopened using Chem3D and an energy minimization was performed. Then, the results were saved as a gic file. In Stage 3, this file was reopened using Gaussian98 and the calculations were performed using the Gaussian® 98 program suite. Gaussian is one of the most widely used quantum chemical program packages for molecular applications, and is used both in industry and in many scientific areas in academia. we have calculated the geometric parameters of the compounds in the ground state using

the Hartree-Fock (HF) (Moller and Plesset, 1934), Becke's three parameter hybrid method (Becke, 1988) with the Lee et al. (1988) correlation functional methods (B3LYP), Becke's exchange functional in combination with the Lee, Yang and Parr correlation functional methods (BLYP) (Lee et al., 1988; Becke, 1993), Becke's three parameter exchange functional combined with gradient corrected correlation functional of Perdew and Wang's 1991 (B3PW91) (Becke, 1993; Predew and Wang, 1992), and 6-31G, 6-31G* and 6-31+G basis set.

The calculation that one asks Gaussian to perform is distributed between many processors to get the answer faster. If one wants to optimize a geometry, it means that one wants Gaussian to adjust the bond lengths, angles and dihedrals to find the lowest energy conformation of the molecule. The command used to tell Gaussian to optimize the molecular geometry is "opt".

The Gaussian program does semi-empirical and *ab initio* calculations. In *ab initio* calculations the important integrals are done directly from first principles. First principles means that the integrals are done either using closed formulas or by doing the integrals numerically. Particularly, *ab initio* method works best for calculating NMR properties (Monajjemi et al., 2010; Monajjemi et al., 2007). Finding a good geometry is called geometry optimization, so "OPT" are used as the keyword.

The calculation will generate an output file called "filename. out". The output file (filename.out) contains a lot of information about the calculation and the results. The content depends on what type of calculation that has been performed and on what print options that was specified. The units are usually Hartree (atomic unit) for energy and Angstrom for distance. There are several different pieces of data that you may need from this. The important information are the Hartree-Fock energy (ΔE), the Mulliken charges, Distance matrix (angstroms), Dipole moment (Debye) and Atomic charge. Distance matrix value is determined (Monajjemi et al., 2008; Mollaamin et al., 2011; Monajjemi et al., 2003).

We used Gaussian 98 at the NMR shift calculation using the HF, B3LYP, BLYP, B3PW91 methods and 6-31G, 6-31G* and 6-31+G basis set. There for "NMR" are used as the keyword (Monajjemi et al., 2003). The calculation will generate an output file called "NMR.out". The output file (NMR.out) contains a lot of information about the NMR shift calculation and are listed in the "GIAO Magnetic shielding tensor (ppm)", such as σ isotropic (ppm) and σ anisotropic (ppm). As the usig the X, Y and z coordinates determinant 3*3 is solved and σ is calculated. Molecular orbital calculations can be used to get good estimates for chemical shifts. In this exercise we calculated the chemical shifts for each of the atom.

In the part of "GIAO magnetic shielding tensor (ppm)" the using the σ lsotropic (ppm), σ anisotropic (ppm) and Eigen values (σ_{11} , σ_{22}, σ_{33}) are calculated parameters such as δ , η and $\Delta\sigma$ (Monajjemi et al., 2006).

We obtained Shielding value for the each atom with Gauss View program (GIAO Magnetic shielding).

A full NBO analysis is obtained in Gaussian when using the POP = NBO keyword. NBO analysis did using the HF, B3LYP, BLYP, B3PW91 methods and 6-31G, 6-31G* and 6-31+G basis set and the output is obtained for each of the molecule (Monajjemi et al., 2006). The main listing of NBOs, displaying the form and occupancy of the complete set of NBOs that span the input AO space and for each orbital gives the type of orbital and the occupancy. We were extracted just BD for 2-center bond and BD* for 2-center antibond from "NBO.output". For example at the Molecule 1.

RESULTS AND DISCUSSION

In this work, we calculated parameters like atomic charges, energy (ΔE), chemical shift anisotropy (δ , asym-

metry parameter (η), chemical shift anisotropy ($\Delta\sigma$), dipole orientation, isotropic, anisotropic, NMR determinant and distance matrix determinant for several benzimidazoles and pyrimidines using the HF method with 6-31G,6-31G* and 6-31+G basis set and B3LYP, BLYP and B3PW91 methods with 6-31G basis set. These parameters are shown in Tables 1 and 2. At present, we considered these parameters.

As shown in Table 1, in Molecule 1, at HF/6-31G, HF/6-31G* and B3PW91/6-31G, N1, N3, C5 atoms contain negative atomic charge values, while in the other levels just N₁ and N₃ atoms contain negative atomic charge values. In Molecule 1 C₂ atom for all methods has the positive value. In Molecule 2, in all of the methods, except BLYP/6-31G level, N₃ and N₅ atoms have the greatest negative atomic charge values and C₄ has the greatest positive amount. In Molecule 3 in all of the methods, N₃ and N₅ atoms have the greatest negative atomic charge values, and C₄ atom has the greatest positive atomic charge amount. In Molecule 4, at all of the methods, N₃ and N₅ have the greatest negative atomic charge, also at all of the methods except HF/6-31+G level C1, N3 and N5 atoms have negative values. In Molecule 5, N₅ has the greatest negative atomic charge. Also at HF/6-31+G, C_6 and C₉ have negative Atomic charge, while in the other levels C_7 and C_{10} have positive atomic charge.

As pointed in Table 1, for Molecule 1, in all of the methods used, except HF/6-31G* chemical shift anisotropy (δ), C₆ has greatest negative value and C₉ is greatest positive value, but at HF/6-31G*, C₆ is greatest negative value and C4 is greatest positive value. For Molecule 2 at all methods, except BLYP, C7 has the greatest positive amount. In HF/6-31G, C₄ has greatest negative amount, while at HF/6-31G* and HF/6-31+G, C2 has greatest negative amount and at B3LYP and B3PW91 methods, C₄ has greatest positive amount. As shown in Molecule 2 for BLYP method has not response. For Molecule 3 in all methods, C₇ has the greatest positive amount. As shown at HF/6-31G, HF/6-31G* and HF/6-31+G, C₂ has greatest negative amount and in B3LYP, BLYP and B3PW91 methods, C₄ has greatest negative amount, also in BLYP method, C2 has greatest positive amount. For Molecule 4 at all methods, N5 has the greatest positive amount and C₄ has greatest negative amount except BLYP method that N₃ has greatest negative amount. For Molecule 5 at all methods, C₄ has the greatest negative amount and N₅ has greatest positive amount.

As pointed in Table 1, in Molecule 1, in all of the methods except BLYP, for the atoms is shown in Table 1, η has the positive amounts. In the BLYP method, N₃ has the negative amount. In Molecule 2, at all of the methods η has positive amounts. Asymmetry parameter (η) value in HF/6-31G ,HF/6-31G^{*}, B3PLY/6-31G and B3PW91/6-31G levels for the C₁ and N₅ atoms, also in HF/6-31+G level for the C₄ and N₅ atoms is the greatest value. In Molecule 3, at all of the methods used for the atoms are shown in Table 1, η has the positive amounts. The η value in all of

Me	thod									ł	łF											B3I	ЛР					BLY	P				B3	PW91			
Bas	is set			6-3	1G					6-3	1G*					6-31	+G											6-3	1G								
Name	Atoms Parameters	A tomic charge	AE(kcal/mol)	õ	η	Δσ	Dinale mament	Atomic charge	AE(kcal/mol)	ð	η	Δσ	Dipole moment	A tomic charge	AE(kcal/mol)	ð	η	Δσ	Dipole moment	Atomic charge	AE(kcal/mol)	δ	η	Δσ	Dipole moment	A tomic charge	AE(kcaVmol)	ð	η	Δσ	Dipole moment	A tomic charge	AE(kcal/mol)	δ	η	Δσ	Dipole moment
Molecule 1	N(1) C(2) N(3) C(4) C(5) C(6) C(9)	-1.0 1.07 -1.03 0.51 -0.31 0.84 0.5	0	-69.32 -110.23 -79.74 132.91 77.73 -118.79 141.60	0.35 0.41 0.04 0.76 0.75 0.63 0.58	-104 -165.34 -120.06 199.37 116.60 178.18 212.40	2.9077	-0.92 1.04 -0.91 0.46 -0.37 0.86 0.55	-108306.009	-51.01 -87.5 -66.00 126.66 67.53 -95.92 115.61	0.13 0.32 0.23 0.82 0.87 0.60 0.54	-76.52 -131.23 -99.00 189.99 101.29 -143.88 173.41	2.3077	-0.95 0.97 -0.91 0.72 0.95 0.51 0.25	-3.669	-66.50 -107.93 -80.48 132.30 77.2 -116.55 142.62	0.32 0.44 0.08 0.76 0.78 0.64 0.57	-99.75 -161.90 -120.72 198.46 115.8 -174.83 +213.9	3.0763	-0.70 0.69 -0.73 0.35 0.003 0.5 0.28	-2253.442	-73.18 -94.21 -78.96 109.94 76.01 -101.34 119.05	0.23 0.12 0.15 0.58 0.47 0.39 0.55	-109.77 -141.32 -118.44 164.91 142.18 -152.01 178.58	2.6230	-0.61 0.60 -0.64 0.31 0.07 0.41 0.23	-2154,553	-75.14 -89.11 -78.23 102.79 73.58 -96.23 110.83	0.16 4.85 -78.23 0.52 0.41 0.26 0.6	-112.72 -133.67 -117.35 154.19 110.37 -144.35 166.25	2.5577	-0.76 0.74 -0.78 0.37 -0.04 0.53 0.3	-2109.190	-72.80 -93.09 -78.78 109.52 75.87 -100.21 118.84	0.30 0.12 0.09 0.58 0.46 0.39 0.53	-109.20 -139.63 -118.17 164.28 113.81 -150.32 178.27	2.6407
Molecule 2	C(1) C(2) N(3) C(4) N(5) C(6) C(7)	-0.10 0.20 -0.98 1.05 -0.93 0.01 -0.03	0	-97.01 -102.42 -70.60 -103.17 -73.18 21.86 130.49	0.74 0.56 0.16 0.69 0.99 0.03 0.53	-145.51 -153.63 -105.91 -154.76 -109.78 32.79 195.73	4.87.36	-0.09 0.11 -0.85 1.00 -0.84 -0.05 0.08	-142.040	-89.76 -97.76 -55.34 -83.72 69.81 20.50 126.16	0.72 0.57 0.29 0.59 0.70 0.30 0.51	-134.65 -146.65 -83.01 -125.58 104.72 30.75 189.24	3.94.88	0.36 -0.18 -0.86 0.92 -0.82 -0.45 0.49	67.93	-98.09 -101.86 -70.68 -101.47 -74.13 22.83 131.13	0.74 0.56 0.21 0.75 0.95 0.02 0.54	-147.14 -152.80 -106.02 -152.21 -111.19 34.25 196.70	5.0465	0.01 0.10 -0.70 0.67 -0.66 -0.10 0.06	-2266.944	-76.32 -82.11 -67.82 -91.73 70.60 22.51 119.02	0.97 0.72 0.11 0.35 0.88 0.17 0.40	-114.48 -123.16 -101.74 -137.60 105.91 33.77 178.54	3.5370		-2.137.458		-		3.0734	-0.00 0.10 -0.75 0.71 -0.70 -0.13 0.03	-2133.175	-75.95 -81.70 -68.02 -90.55 70.98 21.90 118.40	0.97 0.73 0.09 0.36 0.90 0.20 0.39	-113.92 -122.55 -102.03 -135.82 106.47 32.85 177.60	3.5449
Molecule 3	C(1) C(2) N(3) C(4) N(5) C(6) C(7)	-0.22 0.24 -0.98 1.05 -0.93 0.03 0.32	0	87.45 -113.48 -66.56 -107.23 -79.32 19.70 122.72	0.78 0.83 0.14 0.56 0.80 0.34 0.90	131.18 -170.22 -99.84 160.85 -118.99 29.55 184.09	3.0261	-0.22 0.16 -0.87 1.02 -0.86 -0.02 0.33	428.178	77.96 -109.73 -50.5 -85.94 68.12 19.90 95.61	0.82 0.81 0.02 0.47 0.88 0.22 0.97	116.95 -164.60 -75.79 -128.92 102.18 29.84 143.42	2.6019	0.81 -0.54 -0.86 0.93 -0.83 -0.09 -0.09	-6.358	88.20 -112.17 -66.31 -105.62 -80.73 19.92 122.60	0.73 0.85 0.10 0.61 0.77 0.36 0.89	132.30 -168.26 -99.47 -158.43 -121.09 29.89 183.89	3.0585	0.01 0.13 -0.69 0.68 -0.66 -0.08 0.15	-1689.554	81.65 89.71 -64.04 -94.72 -72.58 22.63 97.25	0.51 0.97 0.17 0.24 0.80 0.17 0.95	122.48 134.57 -96.06 -142.08 -108.88 33.95 145.87	2.2427	0.08 0.10 -0.60 0.60 -0.58 -0.09 0.11	-1614.037	78.99 84.10 -62.37 -90.63 -68.98 24.56 -89.70	0.48 0.92 0.15 0.11 0.80 0.05 0.95	118.48 126.15 -93.55 -135.95 -103.47 36.84 -134.56	2.0934	-0.03 0.13 -0.74 0.72 -0.70 -0.11 0.15	-1581.693	81.38 89.91 -63.79 -93.48 -73.56 21.83 98.06	0.51 0.96 0.23 0.24 0.76 0.21 0.90	122.07 134.87 -95.69 -140.22 -110.34 32.75 147.09	2.2583
Molecule 4	C(1) C(2) N(3) H(10) C(4) N(5) H(11)	0.014 0.34 -0.99 0.39 0.37 -0.53 0.23	0	103.86 107.09 -76.23 7.82 -98.90 277.23 2.84	0.38 0.68 0.42 0.74 0.65 0.71 1.00	155.79 160.64 -114.35 11.74 -148.34 415.85 4.27	3.6771	0.17 0.32 -0.83 0.39 0.30 -0.56 0.22	-98.563	10129 102.68 -71.40 -7.89 -87.94 249.57 -4.49	0.39 0.66 0.37 0.86 0.69 0.57 0.69	151.93 154.02 -107.11 -11.84 -131.91 374.36 -6.74	3.55.26	-0.16 0.76 -0.93 0.48 0.45 -0.36 0.25	-6.827	105.32 107.67 -76.11 7.05 -98.46 273.85 -3.07	0.37 0.67 0.39 0.84 0.69 0.71 0.85	157.98 161.50 -114.17 10.58 -147.69 410.78 -4.61	3.78111	0.04 0.32 -0.75 0.32 0.25 -0.40 0.16	-1531,200	91.17 92.96 -74.82 6.15 -78.67 252.92 -2.90	0.30 0.59 0.41 0.86 0.60 0.71 0.81	136.76 139.45 -112.23 9.88 -118.00 379.38 -4.36	3.3905	0.05 0.31 -0.67 0.29 0.23 -0.36 0.14	-1445.587	86.85 88.57 -74.14 5.64 -73.86 242.1744 -2.82	0.28 0.58 0.36 0.83 0.54 0.71 0.73	130.27 132.85 -110.28 8.46 -110.79 363.26 -4.23	3.5129	0.02 0.32 -0.80 0.34 0.25 -0.42 0.20	-1442.518	90.12 91.58 -74.13 6.44 -77.59 252.02 -2.99	0.30 0.60 0.46 0.83 0.61 0.71 0.71	135.18 137.53 -111.20 9.67 -116.38 378.03 -4.49	3.6411
Molecule 5	C(1) C(2) C(4) N(5) C(6) C(9) H(17)	0.03 0.28 0.37 -0.50 0.50 0.54 0.22	0	91.52 84.84 -90.67 298.63 132.12 136.22 5.77	0.57 0.78 0.85 0.82 0.78 0.68 0.66	137.28 127.26 -136.01 447.94 198.18 204.33 8.65	3.7968	0.10 0.18 0.29 -0.52 0.54 0.56 0.22	-157.013	87.42 81.01 -81.37 269.87 106.21 110.08 5.14	0.65 0.84 0.87 0.67 0.75 0.65 0.89	131.13 121.51 -122.06 404.81 159.32 165.13 7.7159	3.6050	0.03 0.60 0.52 -0.32 -0.03 -0.08 0.25	-9.440	93.02 85.75 -90.91 296.75 133.27 137.23 5.66	0.56 0.75 0.88 0.81 0.75 0.65 0.65	139.54 128.63 -136.37 445.13 199.91 205.85 8.49	3.7972	0.06 0.31 0.7 -0.37 0.31 0.33 0.17	1998.091	79.85 77.74 -73.59 277.29 112.38 114.83 4.37	0.44 0.63 0.79 0.86 0.66 0.61 0.80	119.78 116.61 -110.38 415.93 168.57 172.25 6.56	3.6964	0.07 0.31 0.25 -0.34 0.27 0.28 0.15	16 V 07 61-	75.38 75.60 -69.61 266.22 105.09 106.97 3.85	0.49 0.63 0.74 0.89 0.65 0.61 0.87	113.07 113.40 -104.42 399.33 157.64 160.45 5.77	3.6277	0.05 0.31 0.26 -0.39 0.33 0.35 0.21	ST6781-	78.81 76.42 -72.60 275.94 112.65 114.76 4.72	0.45 0.65 0.80 0.87 0.64 0.59 0.76	118.21 114.63 -108.90 413.91 168.97 172.15 7.08	3.7321

Table 1. Values of parameters like atomic charges, ΔE (kcal/mol), chemical shift (δ), asymmetry parameter (η), chemical shift anisotropy ($\Delta \sigma$), dipole moment for active site of studied moleculars obtained using different methods.

Me	thod	HF												B3LY	Р			BLYP				B3PW	91		
Bas	is set	6-310	5			6-31G	*			6-31+0	J			6-31G											
Name	Atoms Parameters	Isotropic (ppm) 0	o Anisotropy(ppm)	a (ppm)	Distance matrix(\mathbf{A}°)	o Isotropic (ppm)	o Anisotropy(ppm)	a (ppm)	Distance matrix(A°)	o Isotropic (ppm)	o Anisotropy(ppm)	a (ppm)	Distance matrix(\mathbf{A}°)	o Isotropic (ppm)	o Anisotropy(ppm)	a (pm)	Distance matrix(A°)	a Isotropic (ppm)	o Anisotropy(ppm)	a (ppm)	Distance matrix(A°)	o Isotropic (ppm)	o Anisotropy(ppm)	a (ppm)	Distance matrix(A°)
Molecule 1	N(1) C(2) N(3) C(4) C(5) C(6) C(9)	127.30 50.54 137.20 35.53 98.62 33.50 -5.50	70.34 116.90 62.23 199.37 116.60 145.65 212.40	1.51E+06 -6.34E+05 1.81E+06 -2.78E+05 4.96E+05 -6.09E+05 5.62E+05	5.7283e+006	133.70 58.91 142.97 36.01 99.74 43.14 11.30	43.55 86.94 61.33 190 101.30 115.32 173.41	2.09E+06 -2.94E+05 2.39E+06 -3.14E+05 6.03E+05 -3.90E+05 1.51E+05	-4.5616e+006	126.44 48.68 138.07 35.89 98.12 33.75 -6.60	66.19 117.18 65.16 198.46 115.79 143.88 213.93	1.52E+06 -5.88E+05 1.84E+06 -2.72E+06 4.81E+05 -5.77E+05 3.09E+07	-1.8744e+007	95.00 50.43 104.56 38.12 84.29 34.00 -6.44	67.96 79.05 68.47 164.91 114.02 105.76 178.58	3.76E+05 -4.13E+05 5.45E+05 -1.10E+05 2.98E+05 -4.53E+05 3.70E+05	-1.6860e+007	82.64 49.00 91.93 36.74 79.17 32.75 -7.72	65.74 66.86 71.33 154.19 110.37 91.63 166.25	-1.61E+05 -3.49E+05 2.45E+05 -6.98E+04 2.40E+05 -4.02E+05 3.08E+05	-2.2336e+007	99.61 54.24 109.08 42.08 88.05 37.84 -2.97	71.28 78.39 64.93 164.28 113.81 104.92 178.27	4.93E+05 -3.91E+05 6.83E+05 -1.27E+05 3.67E+05 -4.55E+05 3.33E+05	-5.2549e+006
Molecule 2	C(1) C(2) N(3) C(4) N(5) C(6) C(7)	96.40 82.34 178.94 47.06 205.31 168.77 66.20	126.78 120.34 61.57 131.06 109.45 32.79 195.73	-9.36E+03 -3.26E+05 4.98E+06 -4.73E+05 7.58E+06 4.75E+06 -2.4E+05	5.9931e+007	96.56 78.05 179.37 55.87 206.04 162.43 65.03	115.99 115.23 53.93 100.02 104.72 30.75 189.24	1.36E+05 -2.85E+05 5.32E+06 -2.47E+05 8.00E+06 4.23E+06 -2.03E+05	3.8946e+007	95.06 82.25 179.20 45.01 203.20 167.97 64.45	128.29 119.31 64.27 133.77 108.64 34.25 196.70	-5.68E+04 -3.15E+05 4.99E+06 -4.33E+05 7.30E+06 4.67E+06 -2.55E+05	4.2233e+007	89.62 77.56 146.25 49.30 177.14 149.40 61.72	113.06 106.01 56.60 93.49 105.91 33.77 178.54	2.04E+05 -4.11E+04 2.56E+06 -3.72E+05 4.79E+06 3.28E+06 -1.02E+05	4.7266e+007			- - - - -	5.1148e+007	93.16 80.31 150.30 53.03 182.13 152.99 65.09	112.32 106.17 55.70 92.44 106.48 32.85 177.60	2.79E+05 986.54 2.81E+06 -3.51E+05 5.23E+06 3.52E+06 -9.21E+04	4.2722e+007
Molecule 3	C(1) C(2) N(3) C(4) N(5) C(6) C(7)	96.21 63.84 172.95 47.71 203.51 172.19 158.94	131.18 156.32 56.96 126.19 107.59 29.55 184.09	2.95E+05 -5.74E+05 4.52E+06 -5.53E+05 7.24E+06 5.05E+06 -4.13E+04	1.1932e+005	97.76 60.55 175.12 58.15 205.56 165.62 23.73	116.95 149.00 38.97 94.79 102.18 29.84 143.42	4.3E+05 -5.28E+05 5.00E+06 -2.71E+05 7.82E+06 4.49E+06 -1.91E+05	1.1494e+005	94.49 63.27 173.18 46.12 201.41 172.05 8.32	132.30 155.79 54.88 127.99 107.24 29.89 183.89	2.75E+05 -5.46E+05 4.54E+06 -5.17E+05 6.9E+06 5.04E+06 -2.55E+06	1.2022e+005	84.34 59.71 139.43 49.63 173.34 151.97 10.72	122.48 134.57 56.24 88.33 98.27 33.95 145.87	2.44E+05 -2.01E+05 2.21E+06 -4.16E+05 4.38E+06 3.45E+06 -7.59E+04	1.4244e+005	79.98 57.65 126.93 48.83 161.10 143.14 11.29	118.48 126.15 53.84 75.89 93.07 36.84 131.76	2.05E+05 -1.26E+05 1.61E+06 -3.68E+05 3.49E+06 2.87E+06 -1.01E+05	1.6208e+005	88.03 62.68 143.88 53.38 178.10 155.58 146.99	122.07 134.87 59.00 87.49 97.48 32.75 147.09	3.08E+05 -1.89E+05 2.47E+06 -3.95E+05 4.78E+06 3.71E+06 -7.82E+04	l. 3900e+005
Molecule 4	C(1) C(2) N(3) H(10) C(4) N(5) H(11)	62.79 72.08 142.02 26.58 60.69 -19.28 25.28	155.79 160.64 81.65 11.74 122.93 415.85 4.27	-3.98E+04 -1.80E+05 2.11E+06 1.73E+04 -4.03E+05 3.94E+06 1.60E+04	1.0764e+005	63.18 72.50 147.60 25.72 63.55 20.02 24.98	151.93 154.02 73.89 11.03 111.55 374.36 5.71	-3.50E+04 -1.27E+05 2.54E+06 1.54E+04 -2.46E+05 1.63E+06 1.51E+04	1.0547e+005	61.39 71.45 143.86 26.25 59.25 -19.43 25.18	157.98 161.50 79.86 10.58 124.96 410.78 4.28	-4.71E+04 -1.80E+05 2.22E+06 1.69E+04 -3.99E+05 3.91E+06 1.58E+04	1.0858e+005	53.96 65.50 117.10 25.41 59.88 -20.68 25.28	136.76 139.45 79.13 9.22 94.56 379.38 3.96	-1.46E+04 -6.35E+04 9.99E+05 1.55E+04 -1.53E+05 3.15E+06 1.60E+04	1.2674e+005	50.13 62.07 108.35 25.07 58.00 -22.67 25.23	130.27 132.85 75.13 8.46 85.74 363.26 3.66	-1.28E+04 -4.91E+04 7.27E+05 1.50E+04 -1.14E+05 2.90E+06 1.59E+04	1.4258e+005	57.67 69.25 120.56 25.30 62.86 -14.61 25.17	135.18 137.53 81.27 9.67 94.10 378.03 4.30	-1.10E+03 -3.22E+04 1.14E+06 1.52E+04 -1.23E+05 2.77E+05 1.57E+04	1.2440e+005
Molecule 5	C(1) C(2) C(4) N(5) C(6) C(9) H(17)	59.53 70.31 61.56 -39.02 7.18 4.81 30.65	137.28 127.26 125.95 447.94 198.18 204.33 8.65	-4.08E+04 -4.91E+04 -2.60E+05 5.29E+06 1.07E+05 2.57E+05 2.80E+04	6.2399e+005	58.69 70.51 65.12 2.26 21.44 19.71 30.33	131.13 121.51 114.30 404.81 159.32 165.13 7.71	-5.50E+04 -3.52E+04 -1.41E+05 2.56E+06 -7.75E+04 1.17E+08 2.72E+04	6.0210e+005	59.28 70.99 59.91 -37.89 7.42 5.11 30.54	139.54 128.63 128.31 445.13 199.91 205.85 8.49	-5.18E+04 -3.72E+04 -2.69E+05 5.19E+06 1.36E+05 2.83E+05 2.77E+04	6.2687e+005	52.38 61.90 58.67 -44.79 2.21 2.91 29.95	119.78 116.61 99.10 415.93 168.57 172.25 6.56	-2.57E+03 -1.06E+04 -9.56E+04 4.44E+06 1.73E+05 2.05E+05 2.64E+04	7.5968e+005	49.79 58.69 55.91 -48.59 -0.25 1.50 29.68	113.07 113.40 91.12 399.33 157.64 160.45 5.77	-1.21E+04 -1.85E+04 -7.60E+04 4.10E+06 1.65E+05 1.73E+05 2.57E+04	8.7896e+005	55.94 65.38 61.71 -38.82 5.10 5.94 29.82	118.21 114.63 98.27 413.91 168.97 172.15 7.08	1.24E+04 1.67E+04 -6.90E+04 3.94E+06 1.52E+05 1.78E+05 2.59E+04	7.3822e+005

Table 2. Values of parameters like isotropic (σ_{iso}), anisotropic (σ_{aniso}) shielding, NMR determinan and distance matrix determinan for active site of studied moleculars obtained using different methods.

2014 Int. J. Phys. Sci.

the methods, except HF/6-31G^{*}, for the C₂ and C₇ is the greatest value. In HF/6-31G^{*}, N₅ and C₇ have the greatest η value. For Molecule 4, η at all of the methods has the positive value. Also at all of the methods, H₁₀ and H₁₁ atoms are the greatest η values. Finally in Molecule 5, η at all of the methods has the positive value. In HF/6-31G, HF/6-31+G and B3PW91/6-31G levels, the η value for the C₄ and N₅ is the greatest value and for the C₁ is the least value, while at HF/6-31G^{*}, B3LYP/6-31G and BLYP/6-31G the η value for the N₅ and H₁₇ is the greatest value and for the C₁ is the least value and for the C₁ is the least value.

The results from Table 1 indicate that in Molecular 1 at HF/6-31G level, the chemical shift anisotropy ($\Delta\sigma$) value for the N_1 , C_2 and N_3 atoms is negative and for the C_4 , C_5 , C_6 and C_9 atoms is positive. In the other methods the $\Delta \sigma$ value for the N₁, C₂, N₃ and C₆ atoms is negative and for the C_4 , C_5 and C_9 atoms is positive. In Molecule 2 at HF/6-31G and HF/6-31+G levels, $\Delta \sigma$ value for the C₁, C₂ N_3 , C₄ and N₅ atoms is negative and for the C₆ and C₇ atoms is positive. While at HF/6-31G*, B3LYP and B3PW91 $\Delta\sigma$ value for the $C_1,~C_2$, N_3 and $C_4~$ atoms is negative and for the N_5 , C_6 and C_7 atoms is positive. In Molecule 3 in all methods $\Delta \sigma$ value for the N₃ and N₅ atoms is negative and for the C_1 and C_7 atoms is positive. In Molecule 4 at HF/6-31G*, HF/6-31+G, B3LYP, BLYP and B3PW91 methods $\Delta \sigma$ value for the N₃, C₄ and H₁₁ atoms is negative, while in HF/6-31G, for the N_3 and C_4 atoms is negative. In Molecule 5 at all methods $\Delta \sigma$ value for the N_5 atom is negative and other atoms are positive.

Dipole orientation (dipole moment) that reported in Table 1, in all molecules (1, 2, 3, 4, 5) for HF/6-31+G is the greatest value. At all molecules for BLYP/6-31G, the dipole moment value is least, except Molecule 5 that for HF/6-31G*, the dipole moment value is least.

 ΔE (kcal/mol) that reported in Table 1, for all molecules at HF/6-31G is zero. Also in all molecules for B3LYP method, ΔE has most negative value except Molecule 1 that in HF/6-31G has most negative value.

As shown in Table 2, in Molecule 1, in all methods except HF/6-31G, $\sigma_{iso}(\sigma \text{ isotropic (ppm)})$ for C₉ is negative, but for other atoms is positive. In HF/6-31G level, σ_{iso} for all atoms is positive. In Molecule 2, at all methods, σ_{iso} for all atoms is positive, and in BLYP method has not response. In Molecule 3, at all methods, σ_{iso} for all atoms is positive. In Molecules 4 and 5, at all methods, except HF/6-31G^{*}, σ_{iso} for N₅ is negative, but for other atoms is positive. In HF/6-31G^{*} level, σ_{iso} for all atoms is positive. In HF/6-31G^{*} level, σ_{iso} for all atoms is positive.

In all molecule (1, 2, 3, 4, 5), $\sigma_{aniso}(\sigma anisotropic (ppm))$, at all methods, for all atoms has positive value. In Molecule 1 in all methods, C₉ has greatest value, in Molecules 2 and 3 in all methods, C₇ has greatest value and in Molecules 4 and 5 in all methods, N₅ has greatest value.

As pointed in Table 2, distance matrix value in Molecule 1, at B3PW91 method is biggest negative amount. In Molecule 2, in all methods, distance matrix has positive amounts and in HF/6-31G has biggest amount. In Molecules 3, 4 and 5 in all methods, distance matrix has has positive amounts and in B has BLYP methods biggest amount.

Table 3a and b show share of orbitals contribute in the bonds (BD for 2-center bond and BD* for 2-center anti bonding. According to Table 3a, in Molecule 1 for HF/6-31G, in the N₁-C₂ bond, BD= 0.7858 sp^{2.08} + 0.6184sp^{2.02} reported. Polarization coefficients of the N_1 - C_2 bond N_1 = 0.7858 and C_2 =0.6184 reported, that sizes of these coefficients show the importance of the hybrid of N1 in the formation of the bond, while in the N_1 -C₆ bond BD= 0.7948 sp^{1.98}+0.6069sp^{2.20} reported. Polarization coefficients of the N_1 - C_6 bond N_1 = 0.7948 and C_2 =0.6069 reported, that sizes of these coefficients show the importance of the hybrid of N_1 in the formation of the bond. In Molecule 2 for HF/6-31G, in the C_2 -N₃ bond, polarization coefficients of this bond C2= 0.6198 and N₃=0.7848 reported, that sizes of these coefficients show the importance of the hybrid of N₃ in the formation of the bond, while in the N_5 -C₆ bond N_5 = 0.7835 and C₆ =0.6214 reported, that sizes of these coefficients show the importance of the hybrid of N₅ in the formation of this bond. In Molecule 3 for HF/6-31G, in the C_4 -N₅ bond, polarization coefficients of this bond $C_4 = 0.6208$ and $N_5 =$ 0.7840 reported, that sizes of these coefficients show the importance of the hybrid of N₅ in the formation of the bond, while in the C₄-N₃ bond C₄ = 0.6208 and N₃ = 0.7840 reported, that sizes of these coefficients show the importance of the hybrid of N₃ in the formation of this bond. In Molecule 4 for HF/6-31G, in the C_2 -N₃ bond, polarization coefficients of this bond $C_2 = 0.6322$ and $N_3 =$ 0.7748 and for the N₃-C₄ bond polarization coefficients N₃ = 0.7896 and C_2 = 0.6136 reported, that sizes of these coefficients show the importance of the hybrid of N₃ in the formation of the bonds. In Molecule 5 for HF/6-31G, at the C_2 - N_3 bond, polarization coefficients of this bond C_2 = 0.6372 and $N_3 = 0.7707$ and for the N_3 - C_4 bond polarization coefficients $N_3 = 0.7922$ and $C_2 = 0.6102$ reported, that sizes of these coefficients show the importance of the hybrid of N₃ in the formation of these bonds reported. As shown in Table 3a and b, for the whole molecular, in all methods at the C-N bond, a polarization coefficient for N is biggest.

Chemical shifts (σ) for each of the atom are shown in Figure 2. As shown in Figure 2, in Molecule 1 at all methods biggest signals are for atoms O₁₁ and O₇. Figure 3 shows that in Molecule 2 at all methods biggest signal is watched for the atoms N₃, N₅ and O₈, except BLYP/6-31G that is not response. The results from Figure 4 indicate that in Molecule 3 at all methods biggest signal is for atom O_8 (aldhyd). Figure 5 shows diagrams which show chemical shifts for each of the atom at the Molecule 4. According to Figure 5 biggest signals are for atoms N₃ and N₅. At all methods σ value for N₅ is more than N₃, but at HF/6-31G^{*}, σ value for N₃ is more than N₅. As shown in Figure 6, in Molecule 5 at all methods biggest signals are for atoms O_{10} and O_{11} . Figures 7 to 11 compare graphs of chemical shifts for molecule with different methods.

Met	hod				HF		
Basi	s set	6-3	31G	6-31	l+G	6-3	31G*
Name	Method Atoms	Bonding	Antibonding	Bonding	Antibonding	Bonding	Antibonding
	N1 – C2	0.7858 sp ^{2.08} + 0.6184sp ^{2.02}	0.6184 sp ^{2.08} - 0.7858 sp ^{2.02}	0.7840 sp ^{2.16} + 0.6207 sp ^{2.03}	0.6207 sp ^{2.16} - 0.7840 sp ^{2.03}	0.7892 sp ^{2.08} d ^{0.01} + 0.6141 sp ^{2.03} d ^{0.01}	$0.6141 \text{ sp}^{2.08} \text{ d}^{0.01} - 0.7892 \text{ sp}^{2.02} \text{ d}^{0.01}$
	N1 – C6	0.7948 sp ^{1.98} +0.6069sp ^{2.20}	0.6069 sp ^{1.98} - 0.7948 sp ^{2.20}	0.7941 sp ^{2.03} + 0.6078 sp ^{2.21}	0.6078 sp ^{2.03} - 0.7941 sp ^{2.21}	$0.7976 \text{ sp}^{1.98} + 0.6031 \text{ sp}^{2.18} \text{d}^{0.01}$	0.6031 sp ^{1.98} - 0.7976 sp ^{2.18} d ^{0.01}
e 1	N1 – H13	0.8646 sp ^{2.19} + 0.5024 s	0.5024 sp ^{2.19} - 0.8646 s	0.8662 sp ^{2.07} + 0.4998 s	0.4998 sp ^{2.07} - 0.8662 s	0.8606 sp ^{2.21} + 0.5092 s	0.5092 sp ^{2.21} - 0.8606 s
olecul	C4 – C12	0.7236 sp ^{2.02} + 0.6902sp ^{2.82}	0.6902 sp ^{2.02} - 0.7236 sp ^{2.82}	0.7261 sp ^{1.98} + 0.6876 sp ^{2.95}	0.6876 sp ^{1.98} - 0.7261 sp ^{2.95}	0.7220 sp ^{2.03} + 0.6919sp ^{2.77} d ^{0.01}	0.6919 sp ^{2.03} - 0.7220 sp ^{2.77} d ^{0.01}
Ŵ	C2 – O8	0.5741 sp ^{2.54} + 0.8188sp ^{1.62}	0.8188 sp ^{2.54} - 0.5741 sp ^{1.62}	0.5940 sp ^{1.94} + 0.8045 sp ^{1.52}	0.8045 sp ^{1.94} - 0.5940 sp ^{1.52}	0.5659 sp ^{2.59} d ^{0.01} + 0.8244 sp ^{1.53} d ^{0.01}	0.8244 sp ^{2.59} d ^{0.01} - 0.5659 sp ^{1.53} d ^{0.01}
	C2 – N3	0.6198 sp ^{2.60} + 0.7848 sp ^{1.88}	0.7848 sp ^{2.60} - 0.6198 sp ^{1.88}	-	-	0.6194 sp ^{2.58} d ^{0.01} + 0.7851 sp ^{1.88}	0.7851 sp ^{2.58} d ^{0.01} - 0.6194 sp ^{1.88}
	N3 – C4	0.7889 sp ^{1.98} + 0.6145 sp ^{2.06}	0.6145 sp ^{1.98} - 0.7889 sp ^{2.06}	-	-	0.7930 sp ^{1.96} + 0.6092 sp ^{2.06} d ^{0.01}	0.6092 sp ^{1.96} - 0.7930 sp ^{2.06} d ^{0.01}
2	C4 – O8	0.5690 sp ^{4.14} + 0.8224 sp ^{6.08}	0.8224 sp ^{4.14} - 0.5690 sp ^{6.08}	-	-	0.5716 sp ^{3.06} d ^{0.01} + 0.8205 sp ^{3.61} d ^{0.02}	0.8205 sp ^{3.06} d ^{0.01} - 0.5716 sp ^{3.61} d ^{0.02}
scule	N5 – C6	0.7835 sp ^{1.78} + 0.6214 sp ^{3.18}	0.6214 sp ^{1.78} - 0.7835 sp ^{3.18}	-	-	0.7844 sp ^{1.78} + 0.6202 sp ^{3.13} d ^{0.01}	0.6202 sp ^{1.78} - 0.7844 sp ^{3.13} d ^{0.01}
Mole	C1 – C7	0.7085 sp ^{2.52} + 0.7057 sp ^{2.66}	0.7057 sp ^{2.52} - 0.7085 sp ^{2.66}	-	-	0.7079 sp ^{2.54} + 0.7063 sp ^{2.67}	0.7063 sp ^{2.54} - 0.7079 sp ^{2.67}
	N3 – C4	0.7904 sp ^{1.96} + 0.6126 sp ^{2.08}	0.6126 sp ^{1.96} - 0.7904 sp ^{2.08}	0.7885 sp ²⁰⁴ + 0.6150 sp ²⁰⁸	0.6150 sp ^{2.04} - 0.7885 sp ^{2.08}	0.7944 sp ^{1.94} + 0.6074sp ^{2.08} d ^{0.01}	0.6074 sp ^{1.94} - 0.7944 sp ^{2.08} d ^{0.01}
	C4 – N5	0.6208 sp ² + 0.7840 sp ^{2.05}	0.7840 sp ² - 0.6208 sp ^{2.05}	0.6228 sp ^{2.01} + 0.7824 sp ^{2.13}	0.7824 sp ^{2.01} - 0.6228 sp ^{2.13}	0.6153 sp ² d ^{0.01} + 0.7883 sp ^{2.03}	0.7883 sp ² d ^{0.01} - 0.6153 sp ^{2.03}
3	N5 – C6	0.7851 sp ^{1.78} + 0.6194 sp ^{3.20}	0.6194 sp ^{1.78} - 0.7851 sp ^{3.20}	0.7869 sp ^{1.79} + 0.6170 sp ^{3.29}	0.6170 sp ^{1.79} - 0.7869 sp ^{3.29}	0.7861 sp ^{1.78} + 0.6181sp ^{3.15} d ^{0.01}	0.6181 sp ^{1.78} - 0.7861 sp ^{3.15} d ^{0.01}
ecule	C2 – N3	0.6222 sp ^{2.56} + 0.7829 sp ^{1.91}	0.7829 sp ^{2.56} - 0.6222 sp ^{1.91}	0.6201 sp ^{2.60} + 0.7845 sp ^{1.91}	0.62017845 sp ^{2.60} - 0.6201 sp ^{1.91}	0.6213 sp ^{2.55} d ^{0.01} + 0.7835 sp ^{1.91}	0.7835 sp ^{2.55} d ^{0.01} - 0.6213 sp ^{1.91}
Mole	C1 – C6	0.7136 sp ^{1.99} + 0.7005 sp ^{2.65}	0.7005 sp ^{1.99} - 0.7136 sp ^{2.65}	0.7143 sp ^{1.99} + 0.6999 sp ^{2.71}	0.6999 sp ^{1.99} - 0.7143 sp ^{2.71}	0.7138 sp ^{1.98} + 0.7004sp ^{2.62} d ^{0.01}	$0.7004 \text{ sp}^{1.98}$ - 0.7138 sp $^{2.62} \text{ d}^{0.01}$
	C1 – N5	0.6468 sp ^{2.98} + 0.7627 sp ^{3.23}	0.7627 sp ^{2.98} - 0.6468 sp ^{3.23}	-	-	$0.6450 \text{ sp}^{2.96} \text{d}^{0.01} + 0.7642 \text{ sp}^{3.21} \text{ d}^{0.01}$	$0.7642 \text{ sp}^{2.96} \text{d}^{0.01} \text{ - } 0.6450 \text{sp}^{3.21} \text{ d}^{0.01}$
4	C2 – N3	0.6322 sp ^{3.10} + 0.7748 sp ^{2.40}	0.7748 sp ^{3.10} - 0.6322 sp ^{2.40}	-	-	0.6306 sp ^{3.08} + 0.7761 sp ^{2.42}	0.7761 sp ^{3.08} - 0.6306 sp ^{2.42}
ecule	N3 – C4	0.7896 sp ^{2.63} + 0.6136 sp ^{2.69}	0.6136 sp ^{2.63} - 0.7896 sp ^{2.69}	-	-	$0.7910 \text{ sp}^{2.66} \text{d}^{0.01} + 0.6118 \text{ sp}^{2.65} \text{d}^{0.01}$	0.6118 sp ^{2.66} d ^{0.01} - 0.7910 sp ^{2.65} d ^{0.01}
Mole	C1 – C6	0.7147 sp ^{1.47} + 0.6994 sp ^{1.93}	0.6994 sp ^{1.47} - 0.7147 sp ^{1.93}	-	-	0.7140 sp ^{1.48} + 0.7001 sp ^{1.92}	0.7001 sp ^{1.48} - 0.7140 sp ^{1.92}

Table 3a. Relative natural bond orbital (NBO) for several active bond in studied molecules by : HF method with 6-31G,6-31G* and 6-31+G basis set.

Table 3a. Contd.

	C2 – C9	0.7160 sp ^{1.47} + 0.6981 sp ^{1.95}	0.6981 sp ^{1.47} - 0.7160 sp ^{1.95}	-	-	0.7151 sp ^{1.47} + 0.6990 sp ^{1.94}	0.6990 sp ^{1.47} - 0.7151 sp ^{1.94}
	C1 – N5	0.6529 sp ^{2.58} + 0.7574 sp ^{3.27}	0.7574 sp ^{2.58} - 0.6529 sp ^{3.27}	0.6545 sp ^{2.53} + 0.7561 sp ^{3.33}	0.7561 sp ^{2.53} - 0.6545 sp ^{3.33}	$0.6507 \text{ sp}^{2.56} \text{d}^{0.01} + 0.7593 \text{ sp}^{3.23} \text{ d}^{0.01}$	$0.7593 \text{ sp}^{2.56} \text{d}^{0.01} \text{ - } 0.6507 \text{ sp}^{3.23} \text{ d}^{0.01}$
	C2 – N3	0.6372 sp ^{2.74} + 0.7707 sp ^{2.52}	0.7707 sp ^{2.74} - 0.6372 sp ^{2.52}	0.6381 sp ^{2.70} + 0.7700 sp ^{2.59}	0.7700 sp ^{2.70} - 0.6381 sp ^{2.59}	0.6352 sp ^{2.71} d ^{0.01} + 0.7724 sp ^{2.52}	$0.7724 \text{ sp}^{2.71} \text{d}^{0.01}$ - $0.6352 \text{ sp}^{2.52} \text{ d}^{0.01}$
2	C4 – N5	0.6298 sp ^{1.83} + 0.7767 sp ^{1.29}	0.7767 sp ^{1.83} - 0.6298 sp ^{1.29}	0.6330 sp ^{1.82} + 0.7742 sp ^{1.32}	0.7742 sp ^{1.82} - 0.6330 sp ^{1.32}	0.6279 sp ^{1.80} d ^{0.01} + 0.7783 sp ^{1.28} d ^{0.01}	0.7783 sp ^{1.80} d ^{0.01} - 0.6279 sp ^{1.28}
scule	C4 – N5	0.6445 sp + 0.7646 sp ^{99.99}	0.7646 sp - 0.6445 sp ^{99.99}	0.6419 sp + 0.7668 sp ^{99.99}	0.7668 sp - 0.6419 sp ^{99.99}	0.6351 sp + 0.7724 sp ^{99.99} d ^{13.35}	0.7724 sp - 0.6351 sp ^{99.99} d ^{13.35}
Mole	N3 – C4	0.7922 sp ^{2.56} + 0.6102 sp ^{2.74}	0.6102 sp ^{2.56} - 0.7922 sp ^{2.74}	0.7905 sp ^{2.62} + 0.6125 sp ^{2.77}	0.6125 sp ^{2.62} - 0.7905 sp ^{2.77}	$0.7932 \text{ sp}^{2.59} \text{d}^{0.01} + 0.6090 \text{ sp}^{2.69} \text{d}^{0.01}$	$0.6090 \text{ sp}^{2.59} \text{d}^{0.01}$ - 0.7932 $\text{sp}^{2.69} \text{ d}^{0.01}$

Table 3b. Relative natural bond orbital (NBO) for several active bond in studied molecules by: B3LYP, BLYP and B3PW91 methods with 6-31G basis set.

Me	ethod	B3L	YP	BL	YP	B3P	W91
Bas	sis set			(6-31G		
Name	Method	Bonding	Antibonding	Bonding	Antibonding	Bonding	Antibonding
	Atoms						
	N1 – C2	0.7842 sp ^{2.08} + 0.6205 sp ^{2.00}	0.6205 sp ^{2.08} - 0.7842 sp ^{2.00}	0.5875 sp ^{2.08} + 0.8092 sp ^{1.47}	0.6208 sp ^{2.08} - 0.7839 sp ^{1.99}	0.7839 sp ^{2.08} + 0.6208 sp ^{1.99}	0.6208 sp ^{2.08} - 0.7839 sp ^{1.99}
	N1 – C6	0.7923 sp ^{1.99} + 0.6102 sp ^{2.17}	0.6102 sp ^{1.99} - 0.7923 sp ^{2.17}	$0.7919 \text{ sp}^{2.00} + 0.6106 \text{ sp}^{2.16}$	0.6106 sp ^{2.00} - 0.7919 sp ^{2.16}	0.7919 sp ^{2.00} + 0.6106 sp ^{2.16}	0.6106 sp ^{2.00} - 0.7919 sp ^{2.16}
1	N1 – H13	0.8593 sp ^{2.28} + 0.5114 s	0.5114 sp ^{2.28} - 0.8593 s	0.8566 sp ^{2.32} + 0.5159 s	0.5159 sp ^{2.32} - 0.8566 s	0.8566 sp ^{2.32} + 0.5159 s	0.5159 sp ^{2.32} - 0.8566 s
ecular	C4 – C12	0.7177 sp ^{2.05} + 0.6964 sp ^{2.77}	0.6964 sp ^{2.05} - 0.7177 sp ^{2.77}	0.7161 sp ^{2.05} + 0.6980 sp ^{2.75}	0.6980 sp ^{2.05} - 0.7161 sp ^{2.75}	0.7161 sp ^{2.05} + 0.6980 sp ^{2.75}	0.6980 sp ^{2.05} - 0.7161 sp ^{2.75}
Mole	C2 – O8	0.5875 sp ^{2.08} + 0.8092 sp ^{1.47}	0.8092 sp ^{2.08} - 0.5875 sp ^{1.47}	0.5884 sp ^{2.06} + 0.8085 sp ^{1.48}	0.8085 sp ^{2.06} - 0.5884 sp ^{1.48}	0.5884 sp ^{2.06} + 0.8085 sp ^{1.48}	0.8085 sp ^{2.06} - 0.5884 sp ^{1.48}
	C2 – N3	0.6228 sp ^{2.59} + 0.7824 sp ^{1.89}	0.7824 sp ^{2.59} - 0.6228 sp ^{1.89}	0.6231 sp ^{2.59} + 0.7821 sp ^{1.89}	0.7821 sp ^{2.59} - 0.6231 sp ^{1.89}	0.6232 sp ^{2.60} + 0.7821 sp ^{1.90}	0.7821 sp ^{2.60} - 0.6232 sp ^{1.90}

Table 3b. Contd.

	N3 – C4	0.7869 sp ^{1.94} + 0.6170 sp ^{2.04}	0.6170 sp ^{1.94} - 0.7869 sp ^{2.04}	0.7867 sp ^{1.93} + 0.6173 sp ^{2.04}	0.6173 sp ^{1.93} - 0.7867 sp ^{2.04}	0.7870 sp ^{1.95} + 0.6169 sp ^{2.05}	0.6169 sp ^{1.95} - 0.7870 sp ^{2.05}
	C4 – O8	0.5948 sp ^{2.18} + 0.8039 sp ^{2.14}	0.8039 sp ^{2.18} 0.5948 sp ^{2.14}	0.5948 sp ^{2.06} + 0.8039 sp ^{1.87}	0.8039 sp ^{2.06} - 0.5948 sp ^{1.87}	0.5954 sp ^{2.20} + 0.8035 sp ^{2.19}	0.8035 sp ²²⁰ - 0.5954 sp ^{2.19}
scular 2	N5 – C6	0.7783 sp ^{1.81} + 0.6279 sp ^{3.15}	0.6279 sp ^{1.81} - 0.7783 sp ^{3.15}	0.7774 sp ^{1.81} + 0.6290 sp ^{3.15}	0.6290 sp ^{1.81} - 0.7774 sp ^{3.15}	0.7779 sp ^{1.81} + 0.6284 sp ^{3.16}	0.6284 sp ^{1.81} - 0.7779 sp ^{3.16}
Mole	C1 – C7	0.7079 sp ^{2.54} + 0.7063 sp ^{2.69}	0.7063 sp ^{2.54} - 0.7079 sp ^{2.69}	0.7080 sp ^{2.55} + 0.7062 sp ^{2.71}	0.7062 sp ^{2.55} - 0.7080 sp ^{2.71}	0.7079 sp ^{2.54} + 0.7063 sp ^{2.69}	0.7063 sp ^{2.54} - 0.7079 sp ^{2.69}
	N3 – C4	0.7880 sp ^{1.93} + 0.6157 sp ^{2.06}	0.6157 sp ^{1.93} - 0.7880 sp ^{2.06}	0.7877 sp ^{1.92} + 0.6161 sp ^{2.06}	0.6161 sp ^{1.92} - 0.7877 sp ^{2.06}	0.7881 sp ^{1.94} + 0.6156 sp ^{2.06}	0.6156 sp ^{1.94} - 0.7881 sp ^{2.06}
	C4 – N5	0.6225 sp ^{1.98} + 0.7827 sp ^{2.01}	0.7827 sp ^{1.98} - 0.6225 sp ^{2.01}	0.6226 sp ^{1.97} + 0.7825 sp ^{1.99}	0.7825 sp ^{1.97} - 0.6226 sp ^{1.99}	0.6222 sp ^{1.98} + 0.7829 sp ^{2.01}	0.7829 sp ^{1.98} - 0.6222 sp ^{2.01}
	N5 – C6	0.7800 sp ^{1.80} + 0.6258 sp ^{3.18}	0.6258 sp ^{1.80} - 0.7800 sp ^{3.18}	0.7792 sp ^{1.80} + 0.6268 sp ^{3.17}	0.6268 sp ^{1.80} - 0.7792 sp ^{3.17}	0.7797 sp ^{1.81} + 0.6261 sp ^{3.19}	0.6261 sp ^{1.81} - 0.7797 sp ^{3.19}
scular	C2 – N3	0.6236 sp ^{2.57} + 0.7817 sp ^{1.91}	0.7817 sp ^{2.57} - 0.6236 sp ^{1.91}	0.6235 sp ^{2.58} + 0.7819 sp ^{1.91}	0.7819 sp ^{2.58} - 0.6235 sp ^{1.91}	0.6239 sp ^{2.58} + 0.7815 sp ^{1.92}	0.7815 sp ^{2.58} - 0.6239 sp ^{1.92}
Mole	C1 – C6	0.7076 sp ² + 0.7066 sp ^{2.62}	0.7066 sp ² - 0.7076 sp ^{2.62}	0.7060 sp ^{2.01} + 0.7082 sp ^{2.60}	0.7082 sp ^{2.01} - 0.7060 sp ^{2.60}	0.7078 sp ² + 0.7065 sp ^{2.62}	0.7065 sp ² - 0.7078 sp ^{2.62}
	C1 – N5	0.6507 sp ^{3.01} + 0.7593 sp ^{3.49}	0.7593 sp ^{3.01} - 0.6507 sp ^{3.49}	0.6515 sp ^{3.03} + 0.7587 sp ^{3.57}	0.7587 sp ^{3.03} - 0.6515 sp ^{3.57}	0.6512 sp ^{3.01} + 0.7589 sp ^{3.50}	0.7589 sp ^{3.01} - 0.6512 sp ^{3.50}
	C2 – N3	0.6330 sp ^{3.17} + 0.7742 sp ^{2.46}	0.7742 sp ^{3.17} - 0.6330 sp ^{2.46}	0.6333 sp ^{3.19} + 0.7739 sp ^{2.49}	0.7739 sp ^{3.19} - 0.6333 sp ^{2.49}	0.6338 sp ^{3.17} + 0.7735 sp ^{2.48}	0.7735 sp ^{3.17} - 0.6338 sp ^{2.48}
4	N3 – C4	0.7888 sp ^{2.68} + 0.6147 sp ^{2.73}	0.6147 sp ^{2.68} - 0.7888 sp ^{2.73}	0.7886 sp ^{2.70} + 0.6149 sp ^{2.73}	0.6149 sp ^{2.70} - 0.7886 sp ^{2.73}	0.7888 sp ^{2.69} + 0.6146 sp ^{2.74}	0.6146 sp ^{2.69} - 0.7888 sp ^{2.74}
scular	C1 – C6	0.7134 sp ^{1.47} + 0.7008 sp ^{1.92}	0.7008 sp ^{1.47} - 0.7134 sp ^{1.92}	0.7132 sp ^{1.47} + 0.7010 sp ^{1.91}	0.7010 sp ^{1.47} - 0.7132 sp ^{1.91}	0.7136 sp ^{1.47} + 0.7006 sp ^{1.92}	0.7006 sp ^{1.47} - 0.7136 sp ^{1.92}
Mole	C2 – C9	0.7150 sp ^{1.46} + 0.6991 sp ^{1.94}	0.6991 sp ^{1.46} - 0.7150 sp ^{1.94}	0.7149 sp ^{1.45} + 0.6992 sp ^{1.93}	0.6992 sp ^{1.45} - 0.7149 sp ^{1.93}	0.7152 sp ^{1.46} + 0.6990 sp ^{1.94}	0.6990 sp ^{1.46} - 0.7152 sp ^{1.94}
S	C1 – N5	0.6525 sp ^{2.62} + 0.7578 sp ^{3.48}	0.7578 sp ^{2.62} - 0.6525 sp ^{3.48}	0.6525 sp ^{2.62} + 0.7578 sp ^{3.48}	0.7578 sp ^{2.62} - 0.6525 sp ^{3.48}	0.6531 sp ^{2.63} + 0.7573 sp ^{3.50}	0.7573 sp ^{2.63} - 0.6531 sp ^{3.50}
ecular	C2 – N3	0.6341 sp ^{2.81} + 0.7733 sp ^{2.58}	0.7733 sp ^{2.81} - 0.6341 sp ^{2.58}	0.6341 sp ^{2.81} + 0.7733 sp ^{2.58}	0.7733 sp ^{2.81} - 0.6341 sp ^{2.58}	0.6345 sp ^{2.81} + 0.7729 sp ^{2.59}	0.7729 sp ^{2.81} - 0.6345 sp ^{2.59}
Mole	C4 – N5	0.6322 sp ^{1.82} + 0.7748 sp ^{1.32}	0.7748 sp ^{1.82} - 0.6322 sp ^{1.32}	0.6322 sp ^{1.82} + 0.7748 sp ^{1.32}	0.7748 sp ^{1.82} - 0.6322 sp ^{1.32}	0.6327 sp ^{1.82} + 0.7744 sp ^{1.33}	0.7744 sp ^{1.82} - 0.6327 sp ^{1.33}

Table 3b. Contd.

C4 – N5	0.6593 sp + 0.7519 sp ^{99.99}	0.7519 sp - 0.6593 sp ^{99.99}	0.6593 sp + 0.7519 sp ^{99.99}	0.7519 sp - 0.6593 sp ^{99.99}	0.6584 sp + 0.7527 sp ^{99.99}	0.7527 sp - 0.6584 sp ^{99.99}
N3 – C4	0.7909 sp ^{2.64} + 0.6122 sp ^{2.77}	0.6122 sp ^{2.64} - 0.7907 sp ^{2.77}	0.7909 sp ^{2.64} + 0.6122 sp ^{2.77}	0.6122 sp ^{2.64} - 0.7907 sp ^{2.77}	0.7907 sp ^{2.66} + 0.6122 sp ^{2.79}	0.6122 sp ^{2.66} - 0.7907 sp ^{2.79}



Figure 1. Structures of Molecules 1, 2, 3, 4 and 5 shown as chemical structures and energy-minimized structures (ball-and-stick model). (a) Molecule 1. 5-acetyl-6-methylpyrimidine-2,4(1H,3H)- dione. (b) Molecule 2. 5-acetyl-6-methylpyrimidine-2,4(1H,3H)- dione. (c) Molecule 3. 1,2,3,4-tetrahydro-2-oxopyrimidine-5-carbaldehyde. (d) Molecule 4. 1H-benzo[d]imidazole. (5) Molecule 5. 5,6-dihydro-1H-benzo[d]imidazole-4,7-dione.









Molecule 1 (HF/6-31G*)







Molecule 1 (B3LYP/6-31G)

1.50E+08

ь



Molecule 1 (BLYP/6-31G)



Molecule 1



1.00E+08 -5.00E+07 -0.00E+00 --5.00E+07 -1 3 5 7 9 111315171921

Atoms

Molecule 1 (B3PW91/6-31G)





Molecule 2 (HF/6-31+G)







Molecule 2 (B3LYP/6-31G)



Molecule 2 (B3LYP/6-31G)

Figure 3. Diagrams which show chemical shifts for each of the atom at the Molecule (2).



Figure 4. Diagrams which show chemical shifts for each of the atom at the Molecule (3).





Molecule 4

Figure 5. Diagrams which show chemical shifts for each of the atom at the Molecule (4).



Molecule 5

Figure 6. Diagrams which show chemical shifts for each of the atom at the Molecule (5).



Figure 7. The graphs of chemical shifts for Molecule 1. (Series1) HF/6-31 g , (Series 2) HF/6-31+g (Series 3) HF/6-31 g*,(Series4) B3LYP/6-31 g , (Series 5) BLYP/6-31 g , (Series 6) B3PW91/6-31 g.



Figure 8. The graphs of chemical shifts for Molecule 2.(Series 1) HF/6-31g , (Series 2) HF/6-31+g (Series 3) HF/6-31 g* , (Series 4) B3LYP/6-31 g , (Series 5) B3PW91/6-31 g.



Figure 9. The graphs of chemical shifts for Molecule 3.(Series 1) HF/6-31 g , (Series 2) HF/6-31+g (Series 3) HF/6-31 g^{*}, (Series 4) B3LYP/6-31 g , (Series 5) BLYP/6-31g , (Series 6) B3PW91/6-31 g.



Figure 10. The graphs of chemical shifts for Molecule 4.(Series 1) HF/6-31 g , (Series 2) HF/6-31+g (Series 3) HF/6-31 g^{*}, (Series 4) B3LYP/6-31 g , (Series 5) BLYP/6-31 g , (Series 6) B3PW91/6-31 g.

	Method			HF				B3LY	(P	BLY	(P	B3PV	V91
	Basis set	6-31	G	6-310	G*	6-31-	+G			6-31	G		
Name	Parameters Atoms	Shielding (ppm)	Degeneracy	Shielding (ppm)	Degeneracy	Shielding (ppm)	Degeneracy	Shielding (ppm)	Degeneracy	Shieldin (ppm)	Degeneracy	Shielding (ppm)	Degeneracy
	N(1)	127.7	1	-	-	-	-	-	-	-	-	-	-
	N(3)	137.4	1	-	-	-	-	-	-	-	-	-	-
	H(13)	26.95	1	-	-	-	-	-	-	-	-	-	-
	H(14)	27.72	1	-	-	-	-	-	-	-	-	-	-
	C(2)	50.7	1	-	-	-	-	-	-	-	-	-	-
	C(4)	35.5	1	-	-	-	-	-	-	-	-	-	-
e 1	C(6)	33.5	1	-	-	-	-	-	-	-	-	-	-
lecul	C(9)	-5.5	1	-	-	-	-	-	-	-	-	-	-
Mo	O(8)	-2.3	1	-	-	-	-	-	-	-	-	-	-
	N(3)	179	1	179.5	1	179.2	1	146.2	1	-	-	150.3	1
	N(5)	205.3	1	206.1	1	203.2	1	177.2	1	-	-	182.2	1
	H(15)	29.26	1	28.78	1	28.81	1	28.31	1	-	-	28.15	1
	H(16)	30.7	1	30.09	1	29.7	1	29.77	1	-	-	29.67	1
	H(17)	29.31	2	28.73	1	29.07	1	28.13	1	-	-	28.07	1
	C(1)	96.3	1	96.54	1	95	1	89.57	1	-	-	93.2	1
e 2	C(2)	82.3	1	78.06	1	82.2	1	77.56	1	-	-	80.31	1
lecule	C(4)	47.06	1	55.87	1	45.02	1	49.3	1	-	-	53.03	1
Mol	O(8)	34.8	1	84.77	1	34.8	1	4.45	1	-	-	10.17	1

Table 4. Ralative GIAO Magnetic shielding for active site of studied moleculars obtained using different methods.

Table 4. Contd.

	N(3)	173	1	175	1	173.2	1	139.4	1	126.96	1	143.89	1
	N(5)	203.5	1	205.5	1	201.4	1	173.34	1	161.11	1	178.1	1
	H(11)	28.94	1	28.57	1	28.39	1	28.06	1	27.82	1	27.92	1
	H(12)	30.94	1	29.93	1	29.65	1	29.47	1	29.18	1	29.38	1
	H(15)	23.17	1	23.76	1	23.08	1	22.8	1	22.63	1	22.71	1
	C(4)	46.8	1	58.1	1	46.1	1	49.64	1	48.83	1	53.39	1
le 3	C(6)	172.2	1	165.61	1	172	1	151.99	1	143.13	1	155.58	1
lecul	C(7)	9	1	27	1	8.6	1	10.73	1	11.3	1	13.2	1
Ma	O(9)	19	1	75	1	26.58	1	-13.26	1	-23.36	1	-7.59	1
	N(3)	142	1	147.61	1	143.88	1	117.1	1	108.35	1	120.57	1
	N(5)	-19.29	1	20	1	-19.43	1	-20.67	1	-22.68	1	-14.6	1
	H(10)	26.59	1	25.73	1	26.26	1	25.41	1	25.08	1	25.3	1
	H(11)	25.29	1	24.98	1	25.18	1	25.29	1	25.23	1	25.17	1
	H(14)	25.81	1	25.36	1	25.63	1	25.55	2	25.47	2	25.42	2
	H(15)	25.95	2	25.45	2	25.73	2	25.6	3	25.48	3	25.47	3
le 4	C(1)	62.8	1	63.19	1	61.39	1	53.97	1	50.13	1	57.69	1
olecu	C(2)	72.09	1	72.5	1	71.4	1	65.5	1	62.09	1	69.25	1
Й	C(4)	60.7	1	63.55	1	59.25	1	59.89	1	58	1	62.86	1
	N(3)	123.66	1	131.33	1	126.45	1	105.9	1	99.58	1	109.49	1
	N(5)	-39	1	2.26	1	-37.8	1	-44.8	1	-48.6	1	-38.82	1
	H(12)	25.38	1	24.89	1	25.1	1	24.45	1	24.2	1	24.4	1
	H(13)	25.21	1	25.06	1	25.19	1	25.18	1	25.12	1	25.07	1
	C(1)	59.55	1	58.7	1	59.28	1	52.39	1	49.79	1	55.94	1
	C(2)	70.31	1	70.5	1	71	1	61.9	1	58.7	1	65.39	1
le 5	C(4)	61.56	1	65.12	1	59.92	1	58.67	1	55.92	1	61.72	1
olecu	C(6)	83	1	79.67	1	80.15	1	75.99	1	73.79	1	78.63	1
Ŵ	C(9)	94.84	1	93.94	1	95	1	88.81	1	86.1	1	91.5	1



Figure 11. The graphs of chemical shifts for Molecule 5.(Series 1) HF/6-31 g , (Series 2) HF/6-31+g (Series 3) HF/6-31 g^{*}, (Series 4) B3LYP/6-31 g, (Series 5) BLYP/6-31g , (Series 6) B3PW91/6-31 g.

Table 4 shows of GIAO magnetic shielding for some of atoms. As shown in Table 4, in Molecule 1, just HF/6-31G response. In Molecule 1, at HF/6-31G, all atoms degeneracy value is same, and N1, N3 atoms contain positive shielding values, while C9 and O8 atoms contain negative shielding values. In Molecule 2, in all of the methods, except BLYP/6-31G level, N_3 and N_5 atoms have the greatest positive shielding values, at HF/6-31G and HF/6-31+G H₁₅ atom, at HF/6-31G* H₁₇ and at B3LYP/6-31G and B3PW91/6-31G O₈ has the lowest shielding amount. In Molecule 3 in all of the methods, N₃ and N₅ atoms have the greatest positive shielding values. At HF/6-31G, HF/6-31G* and HF/6-31+G, C7 atom has the lowest shielding amount, also at B3LYP/6-31G, BLYP/6-31G and B3PW91/6-31G O₉ has negative shielding value. In Molecule 4, at all of the methods, N₅ has the negative shielding value, also N₃ atom has negative value. In Molecule 5, at all of the methods, except HF/6-31G* level, N₅ has the negative shielding value. At HF/6-31G, N₅ has positive shielding value.

Figure 1 shows the formula structures and the energy minimized 3-D chemical structures of 5-acetyl-6-methylpyrimidine-2,4(1H,3H)-dione (molecule 1), 3,4-dihydro-5-phenylpyrimidin-2(1H)-one (Molecule 2), 1,2,3,4-tetrahydro-2-oxopyrimidine-5-carbaldehyde (Molecule 3), 1H-benzo[d]imidazole (Molecule 4) and 5,6-dihydro-1H-benzo[d]imidazole-4,7-dione (Molecule 6) using ball-and-stick model.

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