

Supplementary Information

Synthesis, spectroscopical studies, DFT, ADME, molecular docking analysis, antimalarial, antimicrobial activities of some *E*-imines and XRD structure of (*E*)-*N*-(2-bromobenzylidene)-2-(trifluoromethyl)benzenamine

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DFT structure of imines-Optimized Structure (1-16)

Molecular ESP evaluation of Schiff bases (1-16)

The HOMO and LUMO of the Schiff bases (1-16)

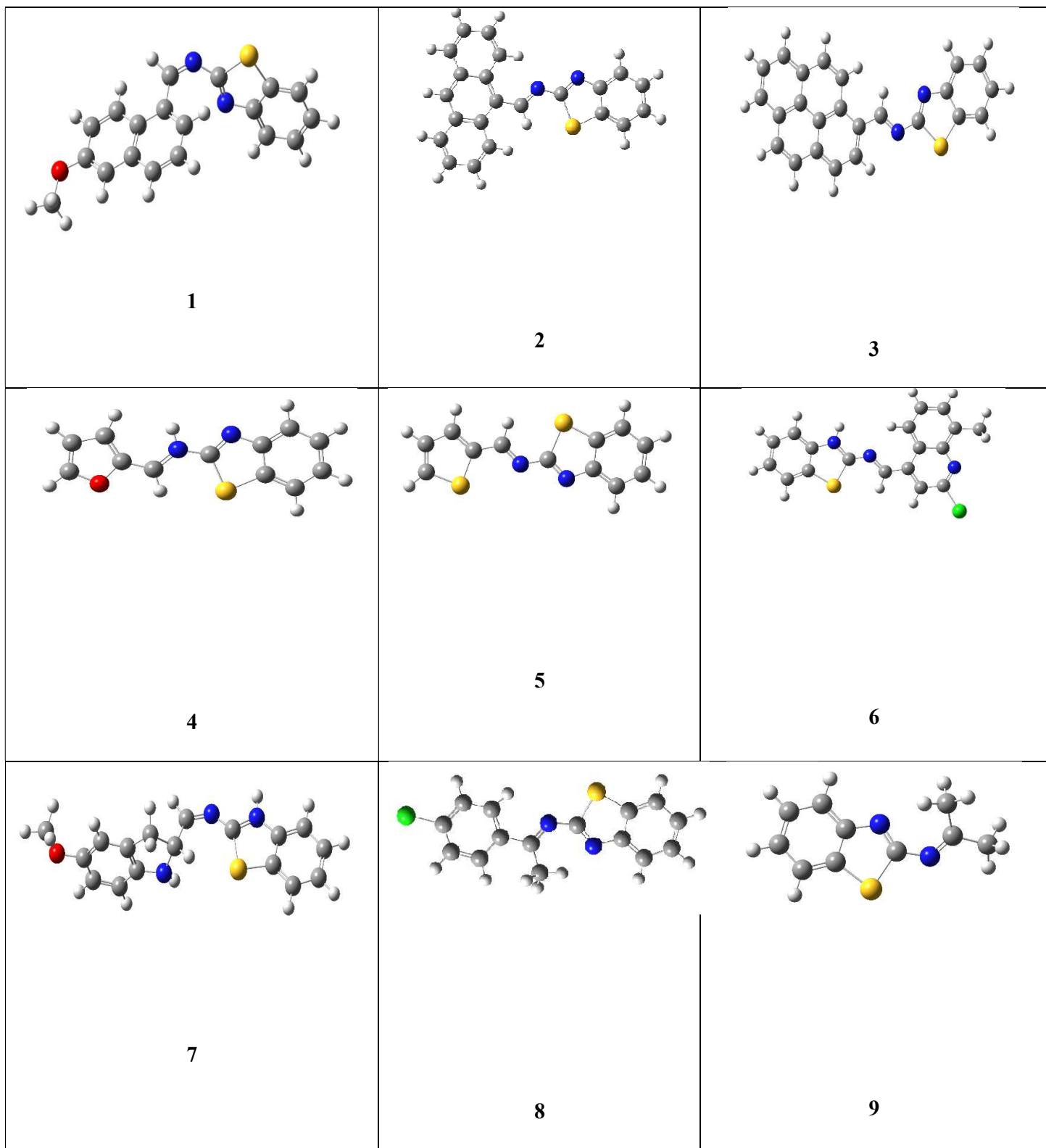
Mulliken Charges of Imines (1-16)

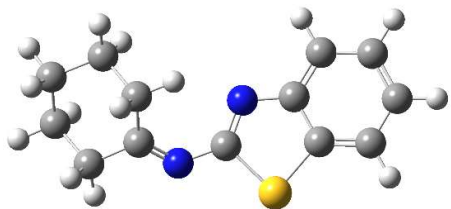
The Molecular Docking 2D and 3D structure of imines (1-16)

The ADME study-Bioavailability Radar for compound (1 to 16)

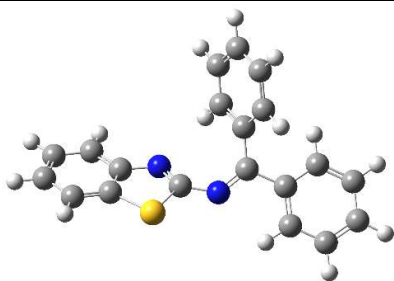
NMR Spectra of *E*-imine compounds (13-16)

DFT structure of imines-Optimized Structure (1-16)

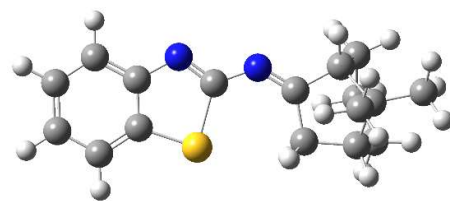




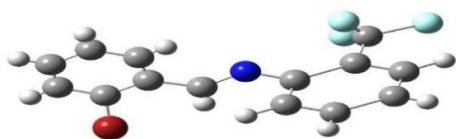
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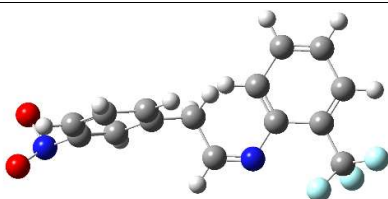
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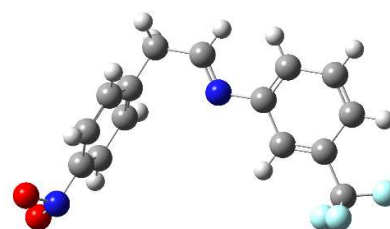
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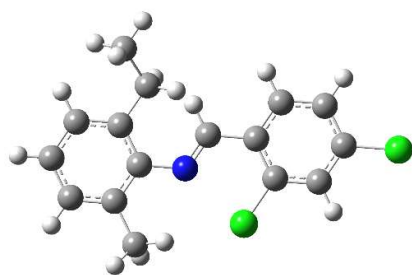
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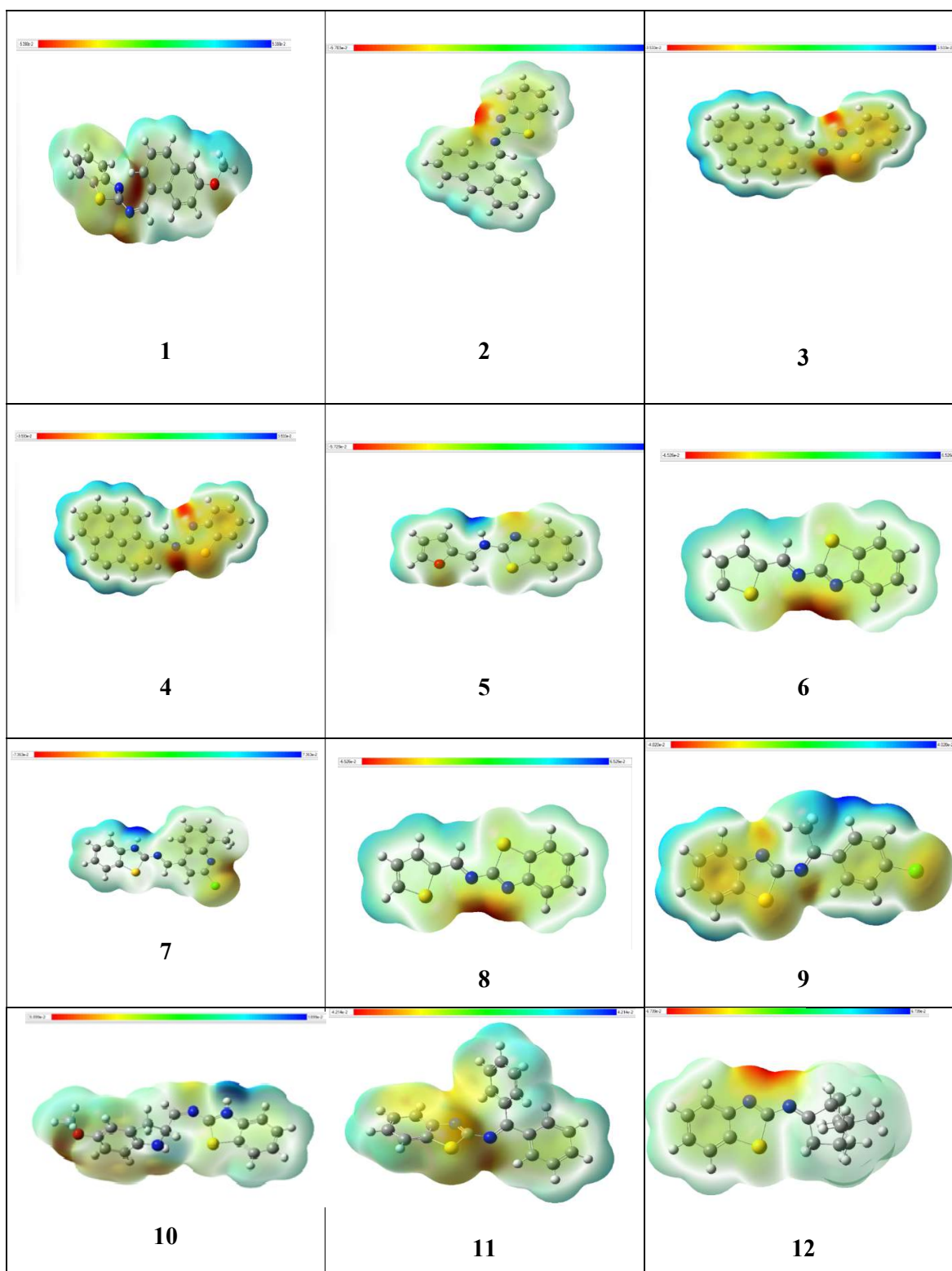


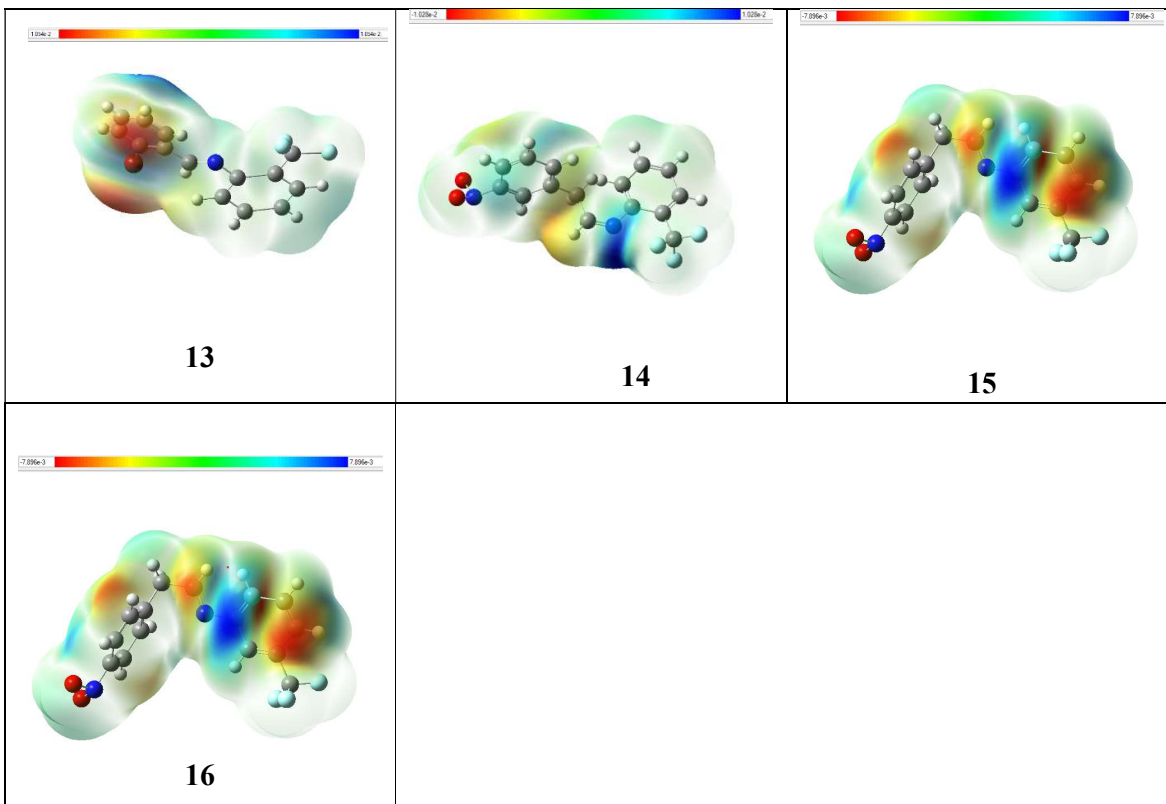
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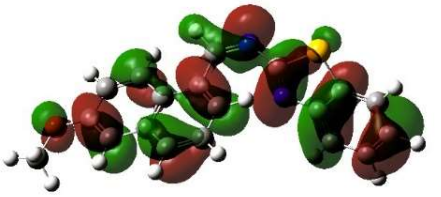

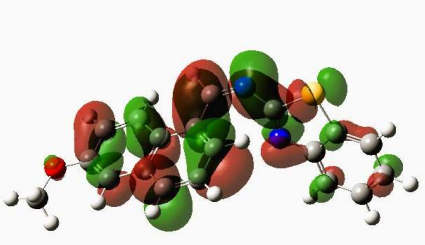
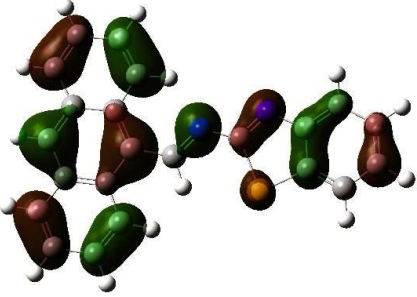

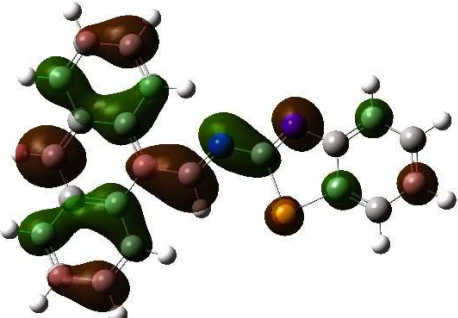
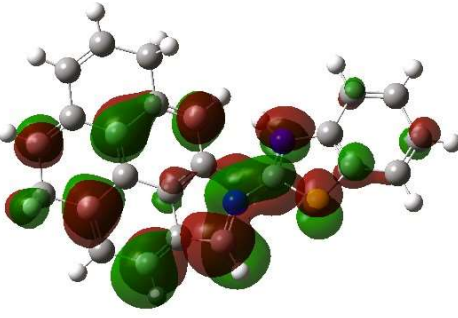

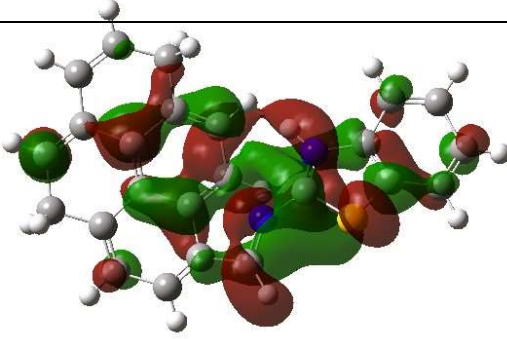
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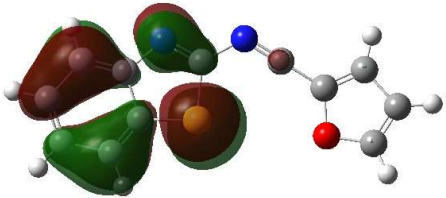

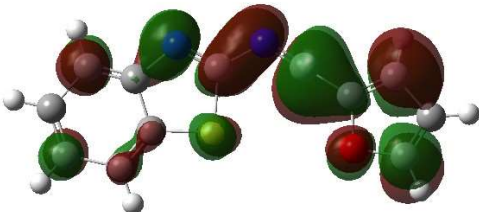
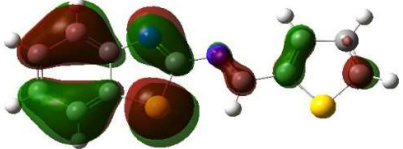
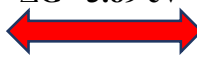
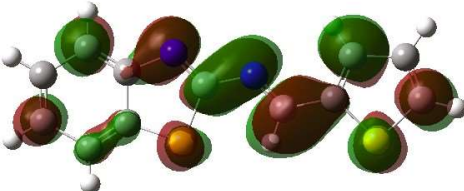
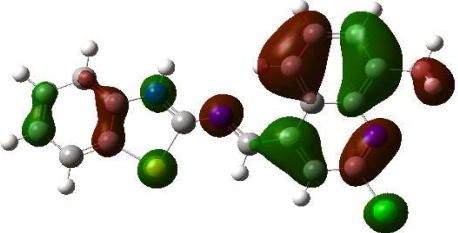

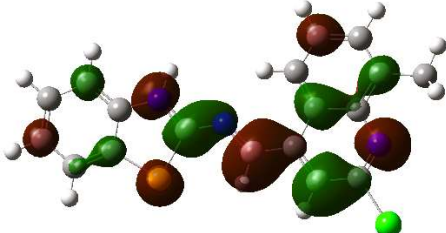
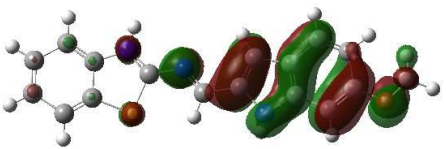

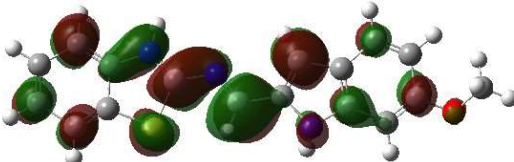
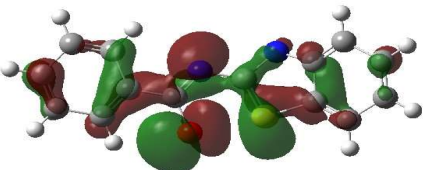

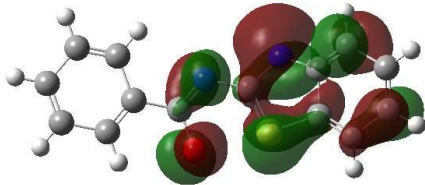
Molecular ESP evaluation of Schiff bases (1-16)

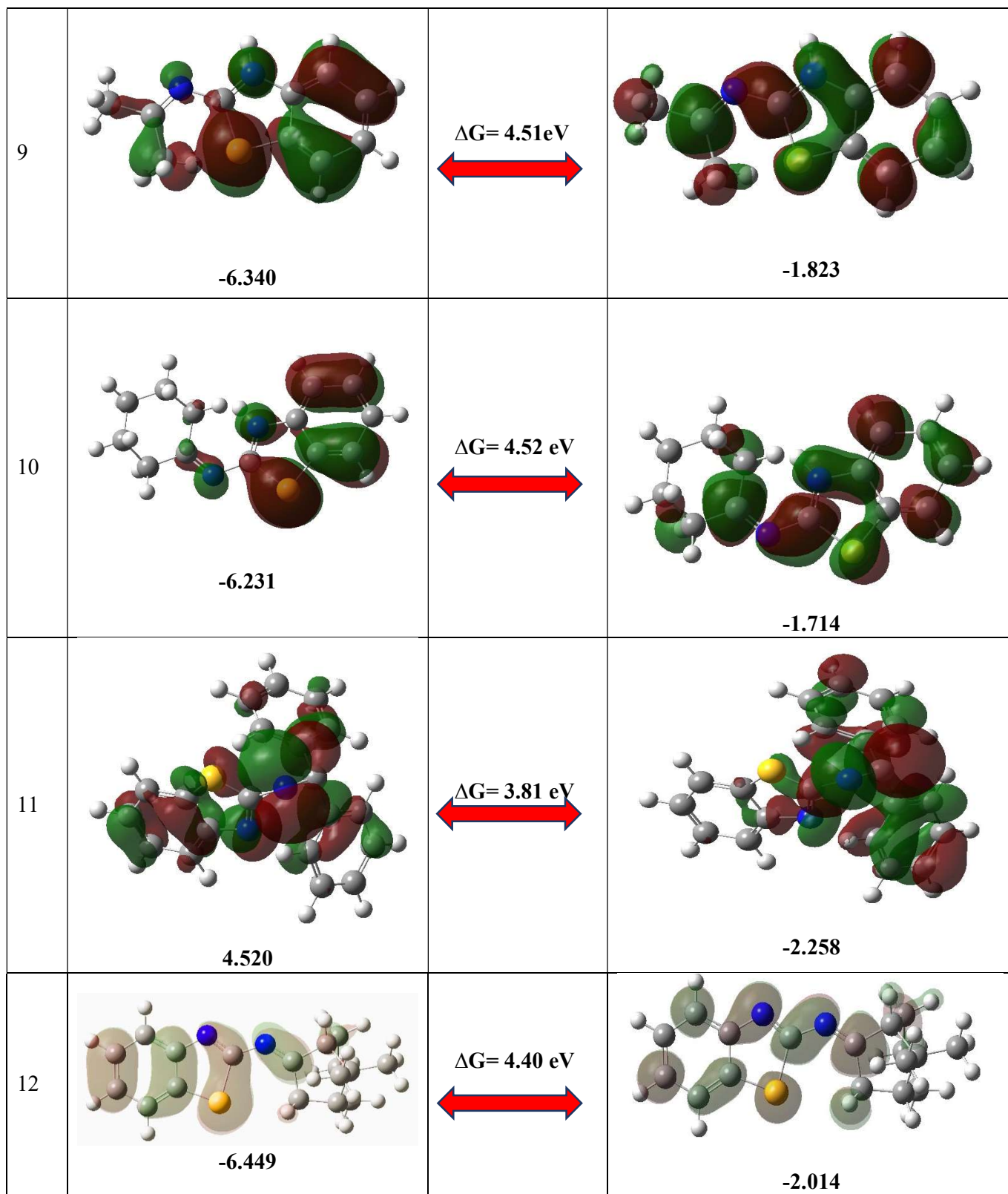


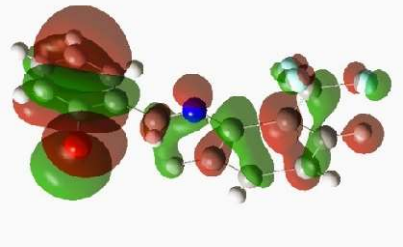
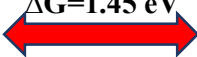
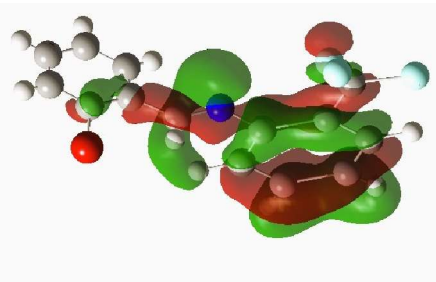
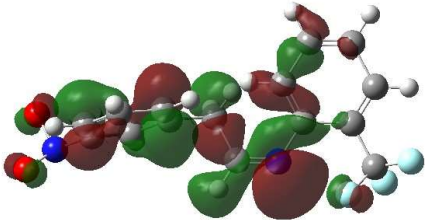

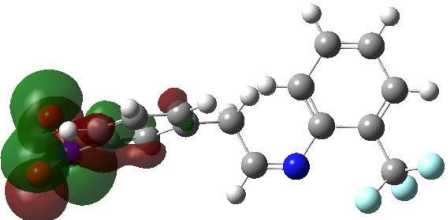
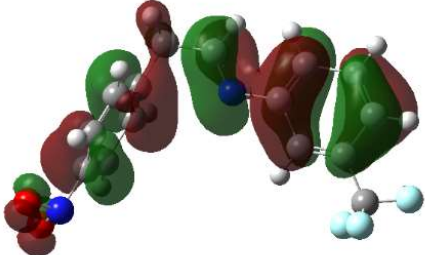

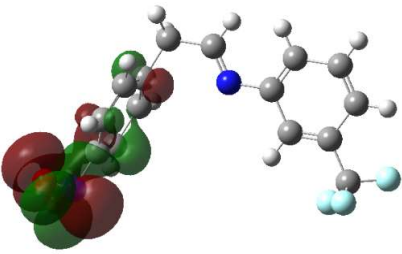
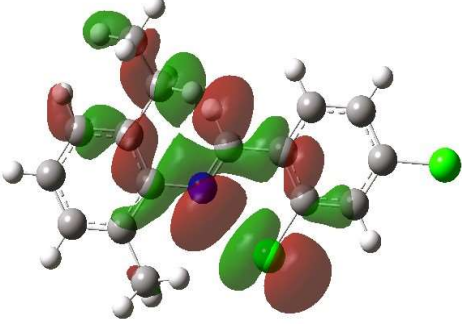
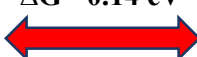
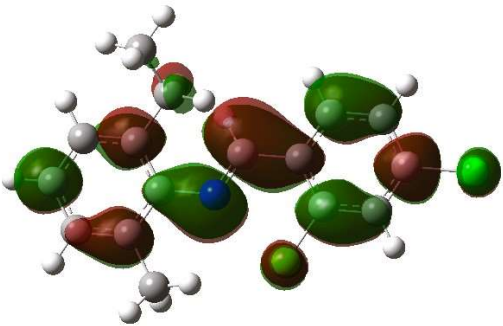


The HOMO and LUMO of the Schiff bases (1-16)

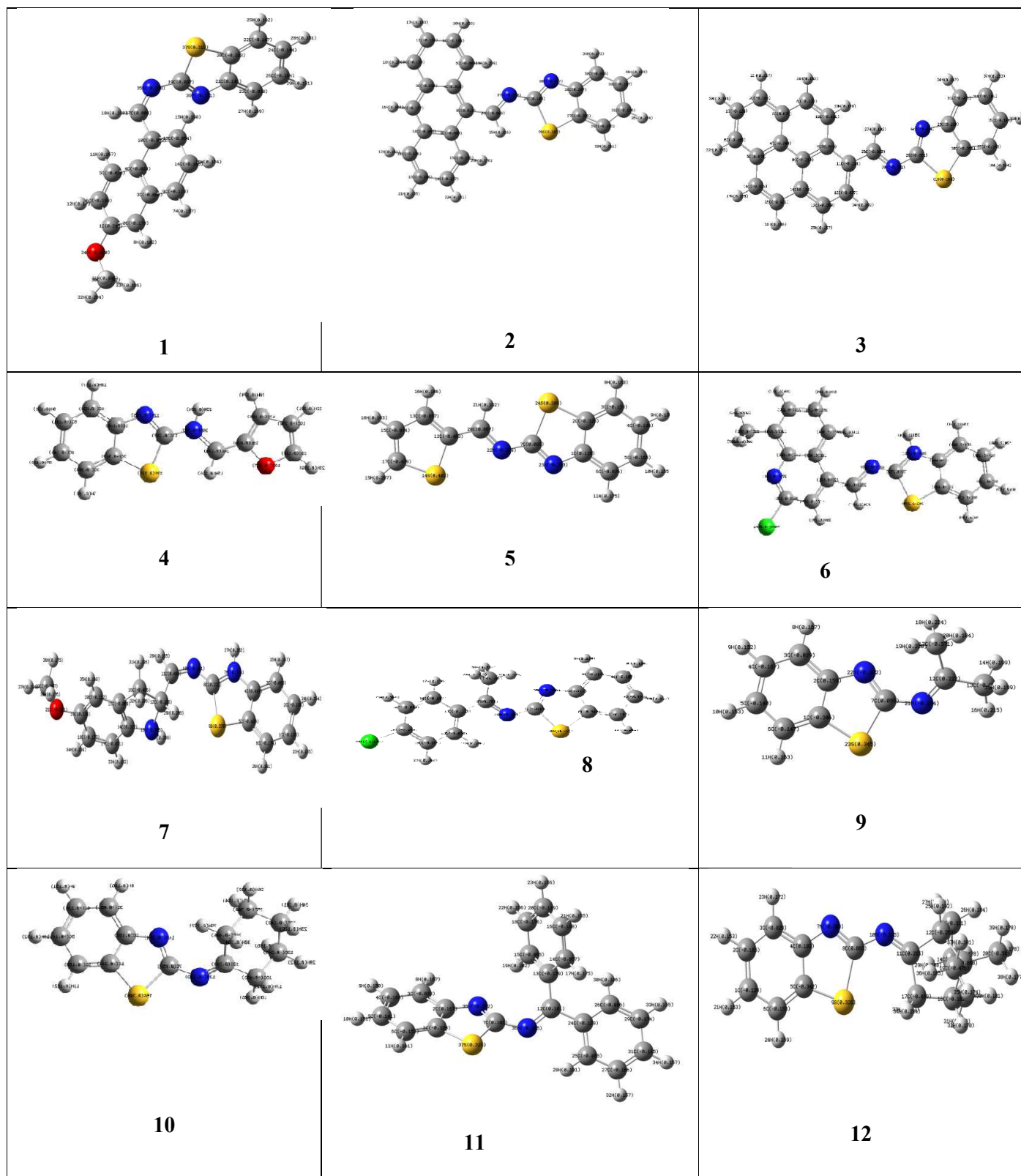
Entry	Homo	Energy gap	Lumo
1	 <p data-bbox="414 714 495 745">-6.068</p>	<p data-bbox="738 577 901 619">$\Delta G = 3.68 \text{ eV}$</p> 	 <p data-bbox="1169 724 1250 756">-2.367</p>
2	 <p data-bbox="414 1218 495 1249">-5.714</p>	<p data-bbox="738 1018 901 1060">$\Delta G = 2.76 \text{ eV}$</p> 	 <p data-bbox="1169 1197 1250 1228">-2.966</p>
3	 <p data-bbox="414 1732 495 1764">-5.741</p>	<p data-bbox="738 1501 901 1543">$\Delta G = 2.96 \text{ eV}$</p> 	 <p data-bbox="1169 1743 1250 1774">-2.775</p>

4	 <p style="text-align: center;">-6.149</p>	<p style="text-align: center;">$\Delta G = 3.77 \text{ eV}$</p> 	 <p style="text-align: center;">-2.394</p>
5	 <p style="text-align: center;">-6.503</p>	<p style="text-align: center;">$\Delta G = 3.89 \text{ eV}$</p> 	 <p style="text-align: center;">-2.612</p>
6	 <p style="text-align: center;">-6.286</p>	<p style="text-align: center;">$\Delta G = 3.19 \text{ eV}$</p> 	 <p style="text-align: center;">-3.102</p>
7	 <p style="text-align: center;">-4.789</p>	<p style="text-align: center;">$\Delta G = 2.69 \text{ eV}$</p> 	 <p style="text-align: center;">-2.095</p>
8	 <p style="text-align: center;">-6.394</p>	<p style="text-align: center;">$\Delta G = 3.80 \text{ eV}$</p> 	 <p style="text-align: center;">2.585</p>

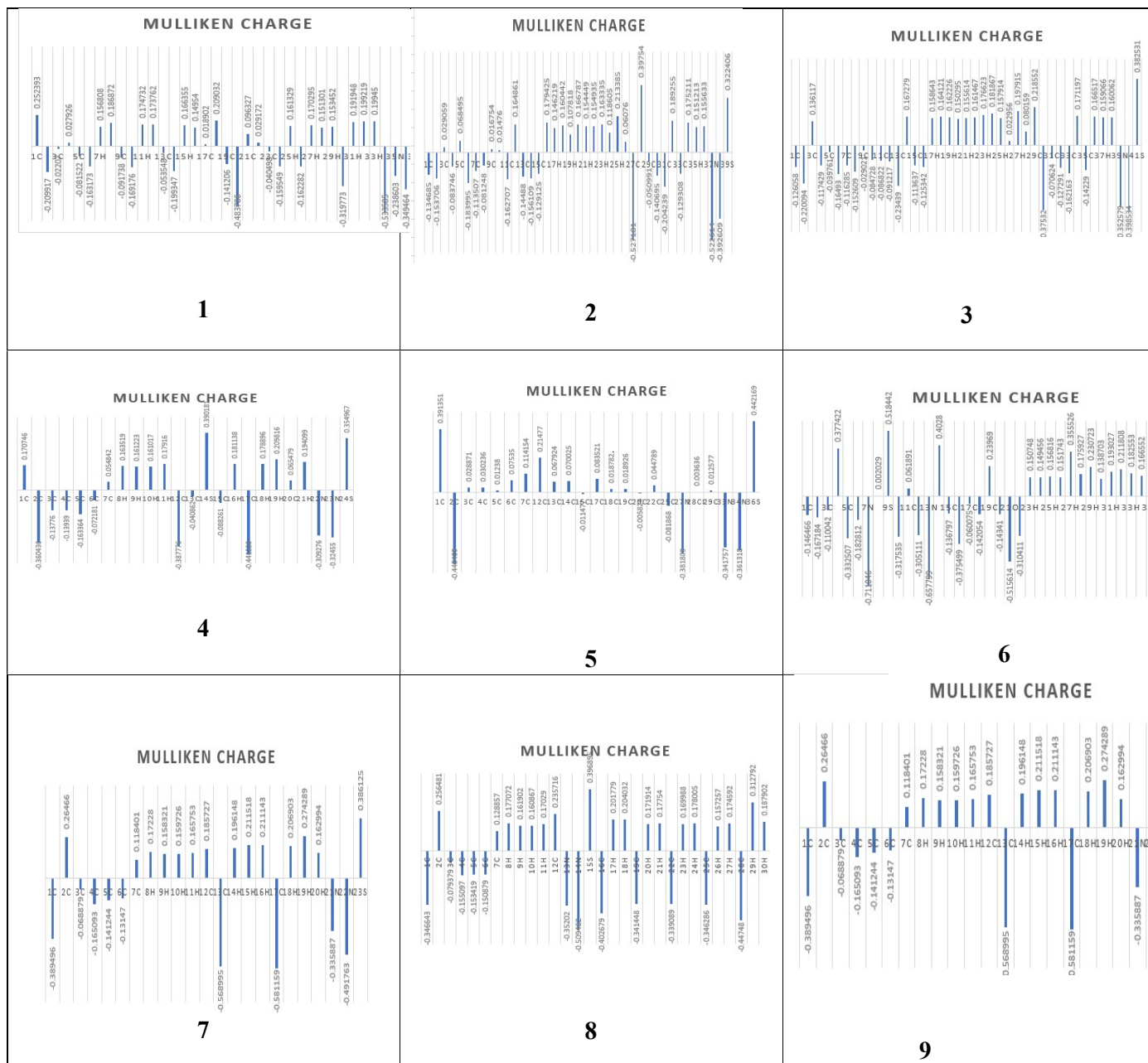


13	 <p style="text-align: center;">-4.843</p>	<p style="text-align: center;">$\Delta G = 1.45 \text{ eV}$</p> 	 <p style="text-align: center;">1.453</p>
14	 <p style="text-align: center;">-7.047</p>	<p style="text-align: center;">$\Delta G = 2.12 \text{ eV}$</p> 	 <p style="text-align: center;">-4.925</p>
15	 <p style="text-align: center;">-7.292</p>	<p style="text-align: center;">$\Delta G = 2.66 \text{ eV}$</p> 	 <p style="text-align: center;">-4.626</p>
16	 <p style="text-align: center;">-5.905</p>	<p style="text-align: center;">$\Delta G = 0.14 \text{ eV}$</p> 	 <p style="text-align: center;">-1.932</p>

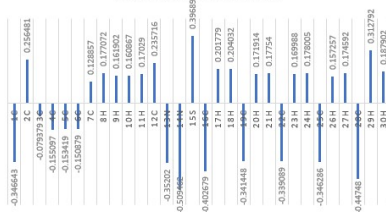
MULLIKEN CHARGES OF Imines 1-16



Mulliken charges of imines (1-16)



MULLIKEN CHARGE



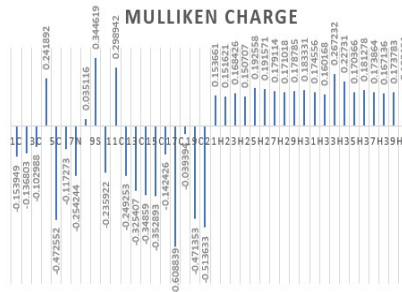
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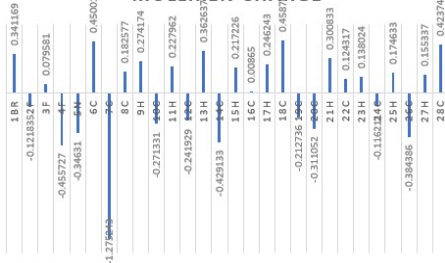
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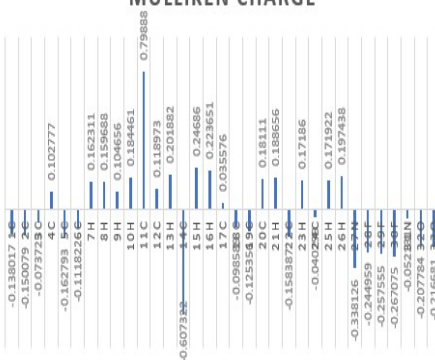
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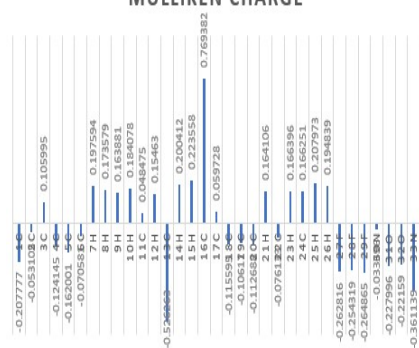
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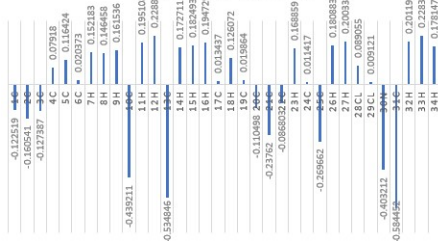
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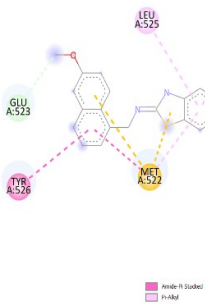
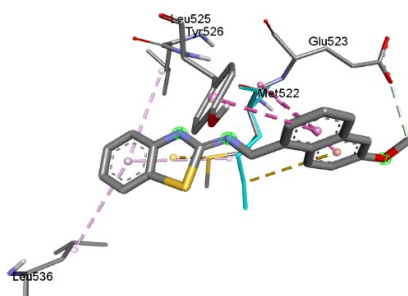
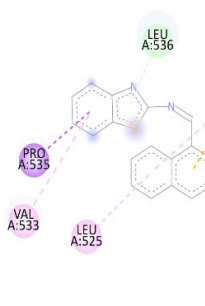
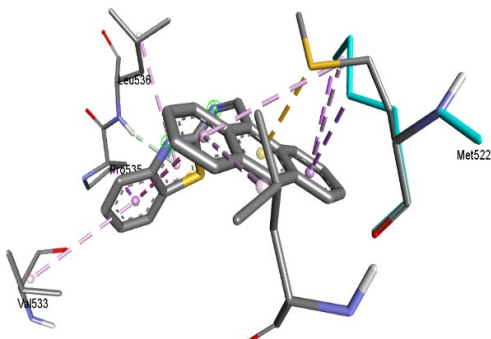
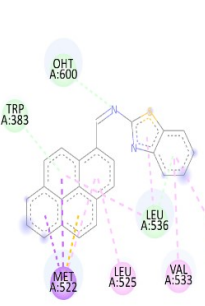
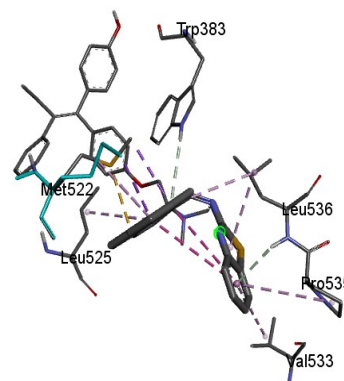
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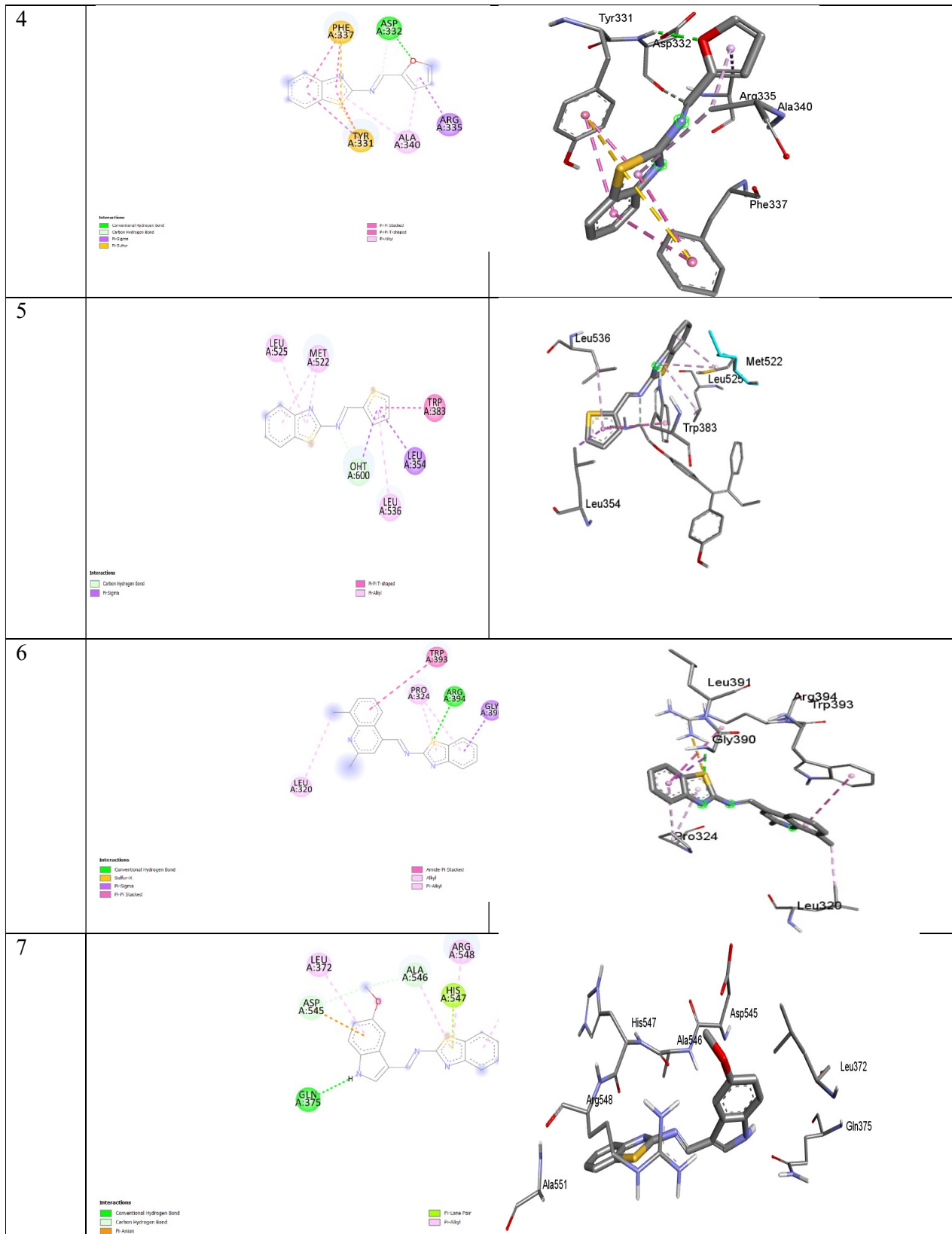
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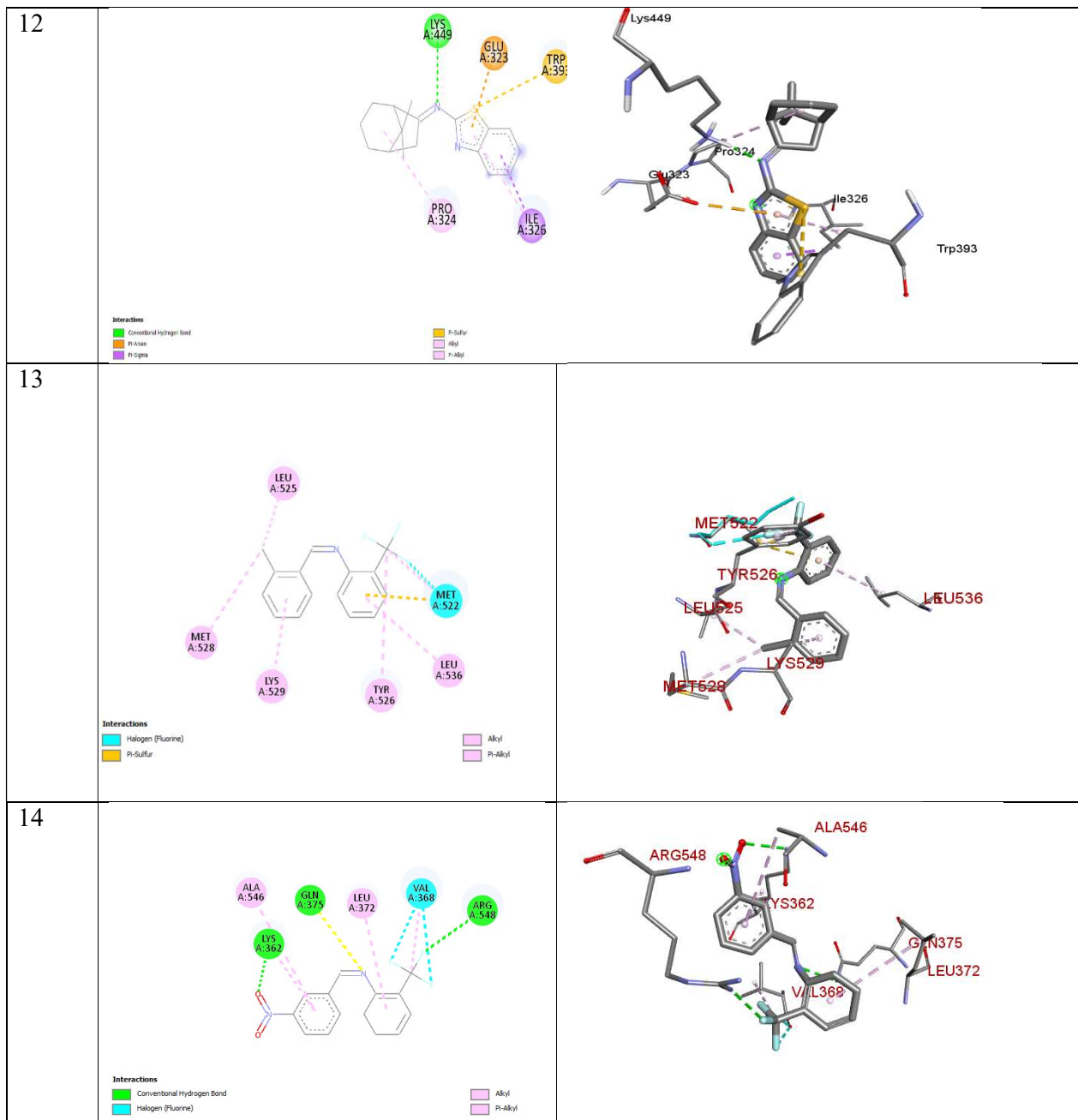
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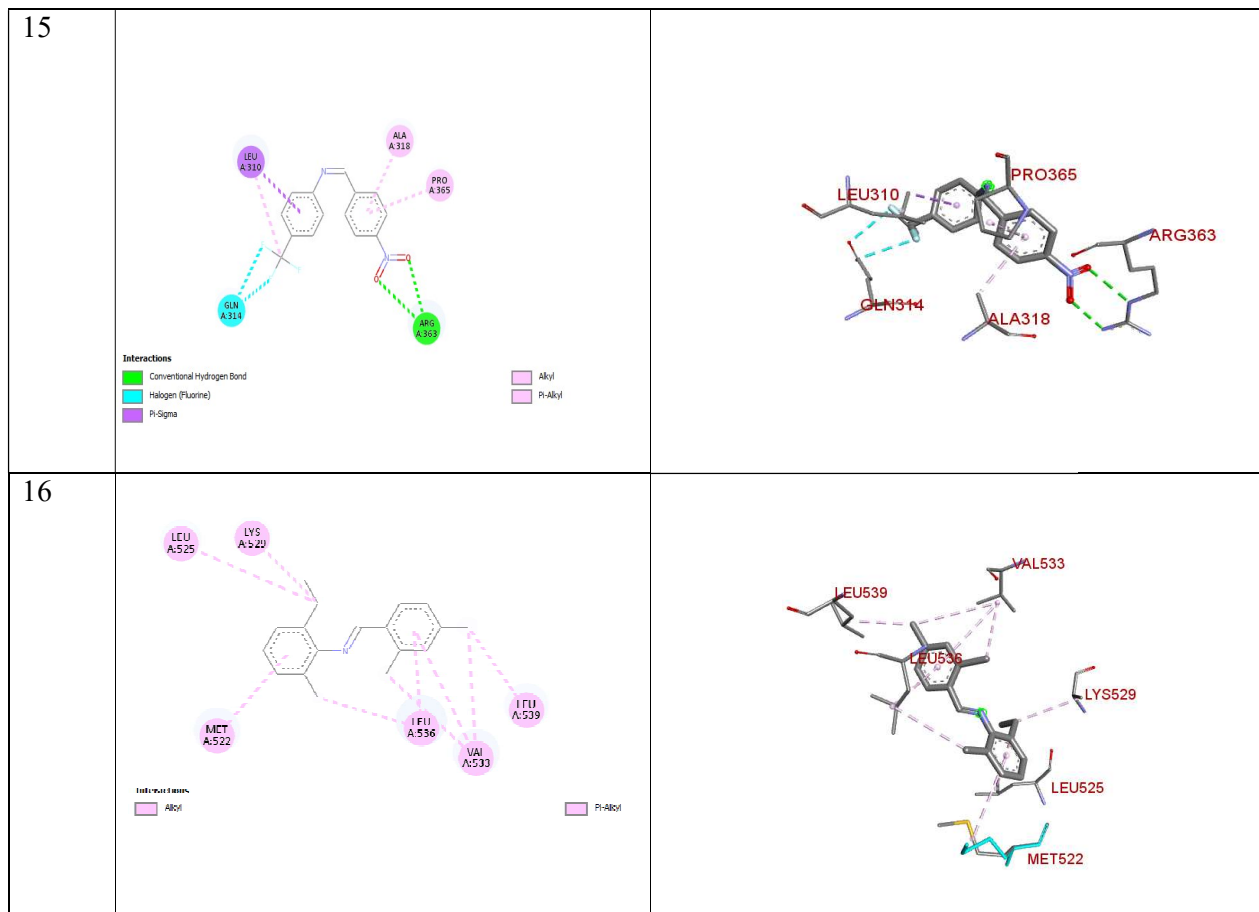
The Molecular Docking 2D and 3D structure of imines (1-16)

Entry	2D	3D
1	 <p>Interactions</p> <ul style="list-style-type: none"> ■ Carbon-Hydrogen bond ■ Pi-Sulfur ■ Pi-Pi stacked ■ Pi-T-shaped ■ Pi-Sigma ■ Pi-Allyl 	
2	 <p>Interactions</p> <ul style="list-style-type: none"> ■ Carbon-Hydrogen bond ■ Pi-Sulfur ■ Pi-Allyl ■ Pi-Sigma 	
3	 <p>Interactions</p> <ul style="list-style-type: none"> ■ Carbon-Hydrogen bond ■ Pi-Donor hydrogen bond ■ Pi-Sulfur ■ Pi-Allyl ■ Pi-Sigma 	


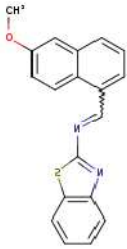
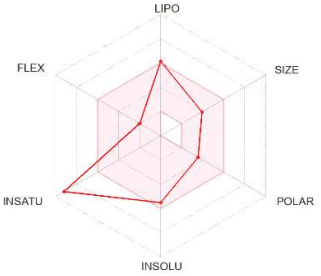

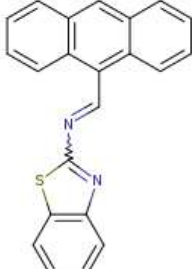

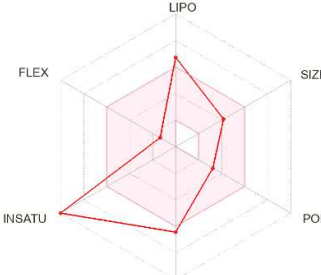

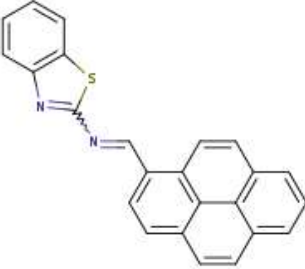
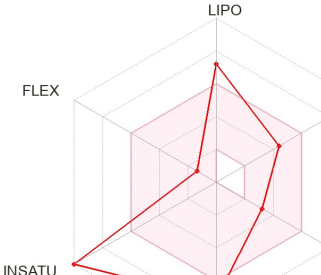


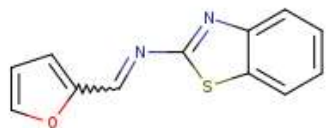
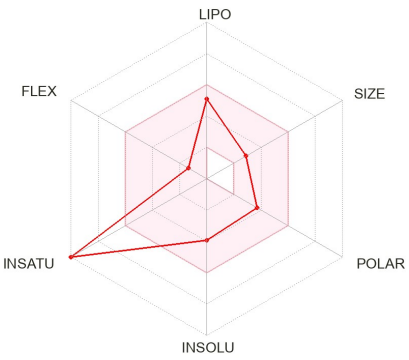
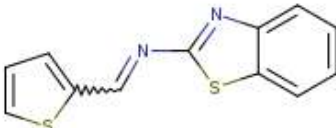
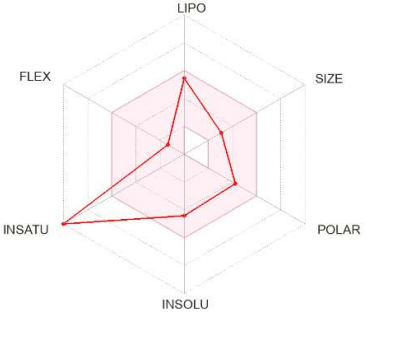
8	<p>Interactions</p> <ul style="list-style-type: none"> Pi-Sulfur Pi-Allyl
9	<p>Interactions</p> <ul style="list-style-type: none"> Sulfur-H Pi-Sulfur Pi-Allyl
10	<p>Interactions</p> <ul style="list-style-type: none"> Cationic-Hydrogen bond Carbon-Hydrogen bond Pi-Amino Pi-Sulfur Allyl Pi-Allyl
11	<p>Interactions</p> <ul style="list-style-type: none"> Pi-Amino Pi-Sulfur Inductor Pi Pi-T shaped Pi-Allyl


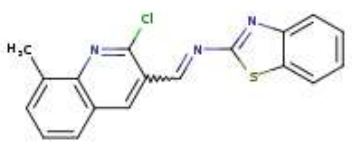
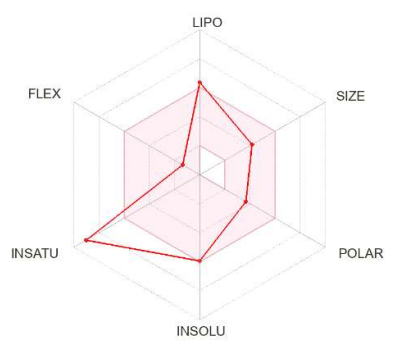

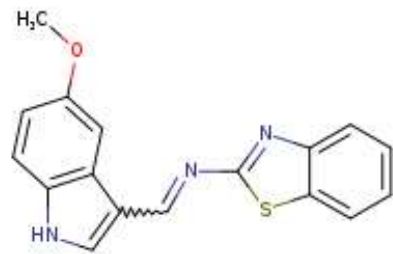
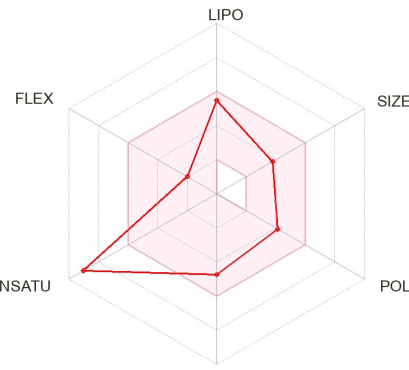

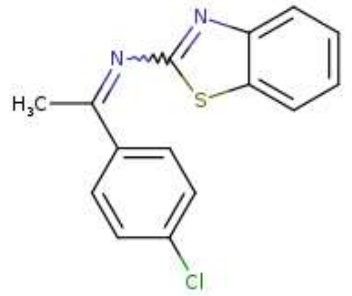




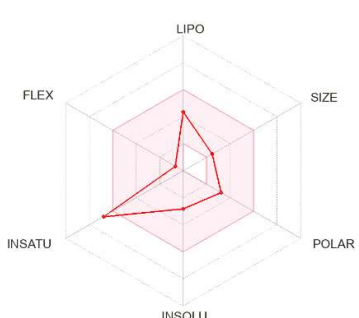
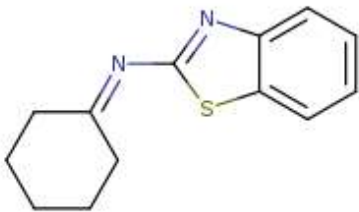
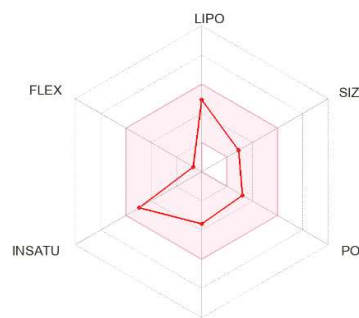


ADME study-Bioavailability Radar for compound (1 to 16)

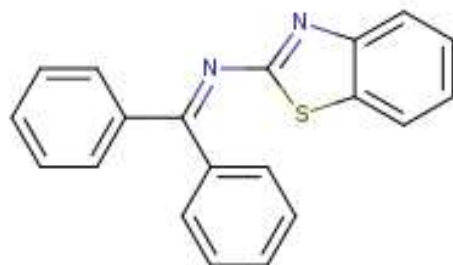
Entry	Compound	Radar
1	<p>  </p>  <p> <chem>COC1=CC=C2C(C=NC3=NC4=C(S3)C=CC=C4)=CC=CC2=C1</chem> </p>	
2	<p>  </p>  <p>  <chem>S1C2=C(C=CC=C2)N=C1\N=C\C1=C2C=CC=CC2=CC2=C1C=CC=C2</chem> </p>	
3	<p>  </p> 	

	<p>S1C2=C(C=CC=C2)N=C1\N=C/C1=CC=C2C=CC3=CC=CC4=C3C2=C1C=C4</p>	
<p>4</p>	<p>⚡ ⊕ ⬡ 🔪 Σ</p>  <p>O1C=CC=C1\C=N/C1=NC2=CC=CC=C2S1</p>	
<p>5</p>	 <p>⚡ ⊕ ⬡ 🔪 Σ</p> <p>S1C=CC=C1C=NC1=NC2=CC=CC=C2S1</p>	

<p>6</p>	<p>  </p>  <p> <chem>CC1=CC=CC2=C1N=C(Cl)C(\C=N\C1=NC3=CC=CC=C3S1)=C2</chem> </p>	
<p>7</p>	<p>  </p>  <p> <chem>COC1=CC=C2C(=C1)NC(=C2)N=C3C=CC=CC=C3S2</chem> </p>	
<p>8</p>	<p>  </p> 	

	<chem>C\C(=N/C1=NC2=CC=CC=C2S1)C1=CC=C(Cl)C=C1</chem>	
9	<p>⚡ ⊕ ⬡ 📄 Σ</p>  <p><chem>CC(C)=NC1=NC2=CC=CC=C2S1</chem></p>	
10	<p>⚡ ⊕ ⬡ 📄 Σ</p>  <p><chem>C1CCC(CC1)=NC1=NC2=CC=CC=C2S1</chem></p>	

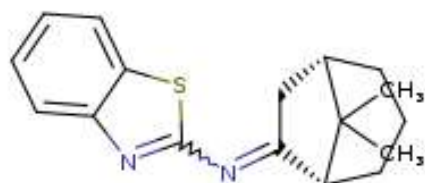
11



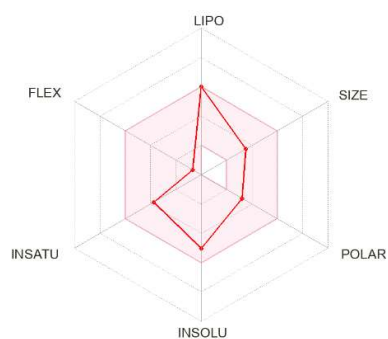
S1C(N=C(C2=CC=CC=C2)C2=CC=CC=C2)=NC2=CC=CC=C2
C12

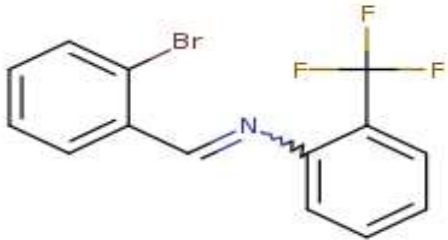
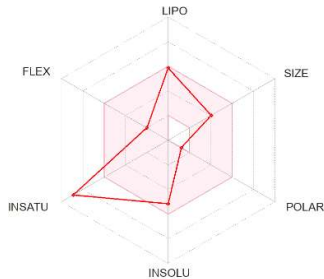
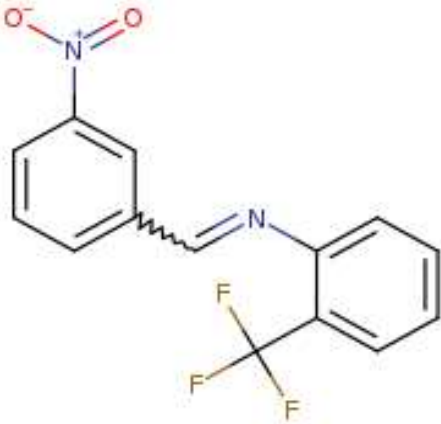
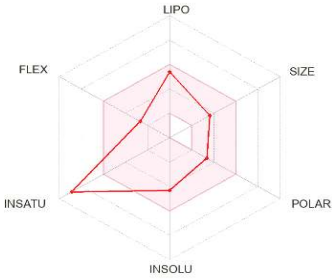
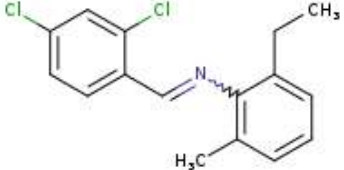
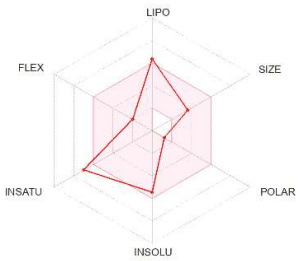



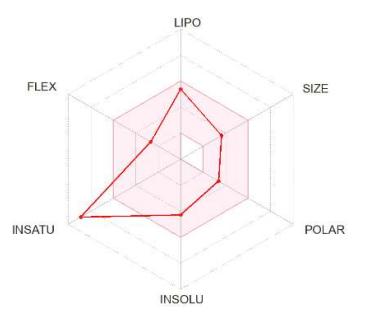
12



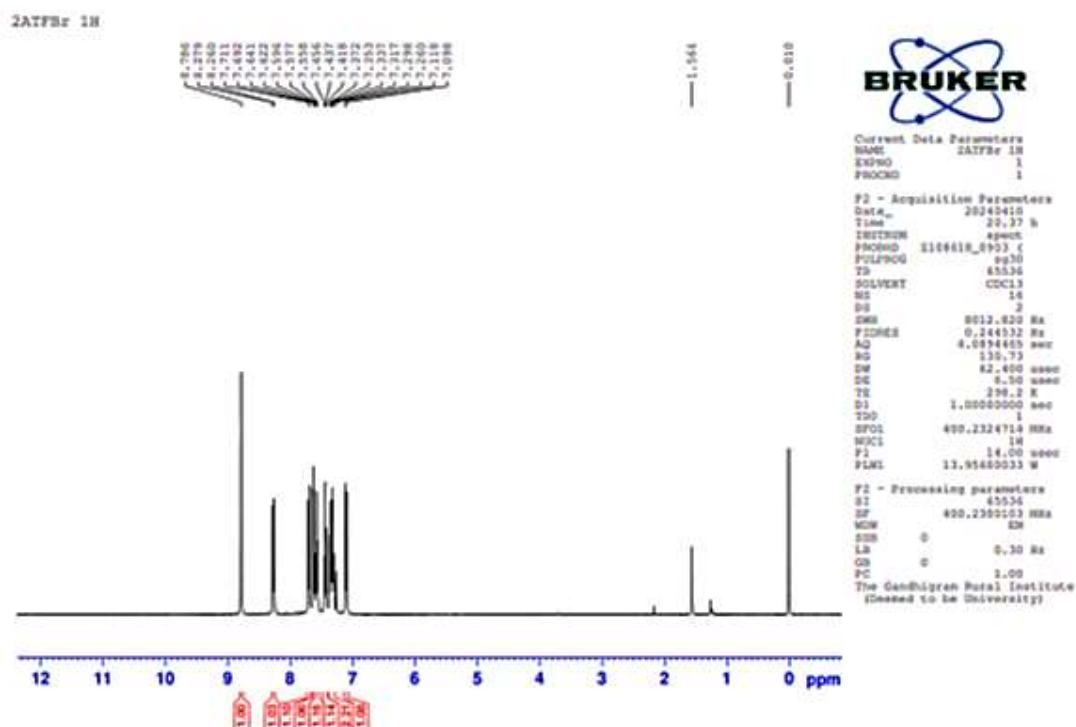
CC1(C)[COOH]2CC(=NC3=NC4=CC=CC=C4S3)[COH]1C
CC2



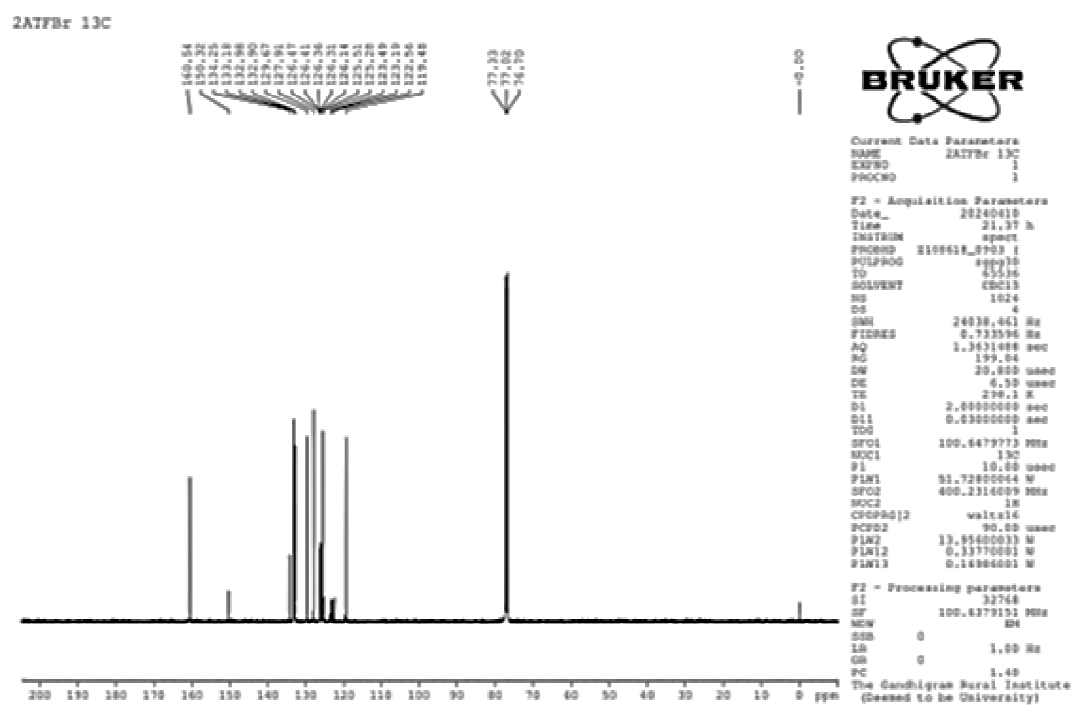
<p>13</p>	<p>⚡ ⊕ ⬡ 🖋 Σ</p>  <p><chem>FC(F)(F)C1=CC=CC=C1\N=C/C1=CC=CC=C1Br</chem></p>	<p>⚡ ⊕ 🖋 Σ ⬡</p> 
<p>14</p>	<p>⚡ ⊕ ⬡ 🖋 Σ</p>  <p><chem>[O][N+](=O)C1=CC(\C=N/C2=CC=CC=C2C(F)(F)F)=CC=C1</chem></p>	
<p>15</p>	<p>⚡ ⊕ ⬡ 🖋 Σ</p> 	

	<chem>CCC1=CC=CC(C)=C1\N=C/C1=CC=C(Cl)C=C1Cl</chem>	
16	 <chem>CCC1=CC=CC(C)=C1\N=C/C1=CC=C(Cl)C=C1Cl</chem>	

NMR Spectra of *E*-imine compounds (13-16)

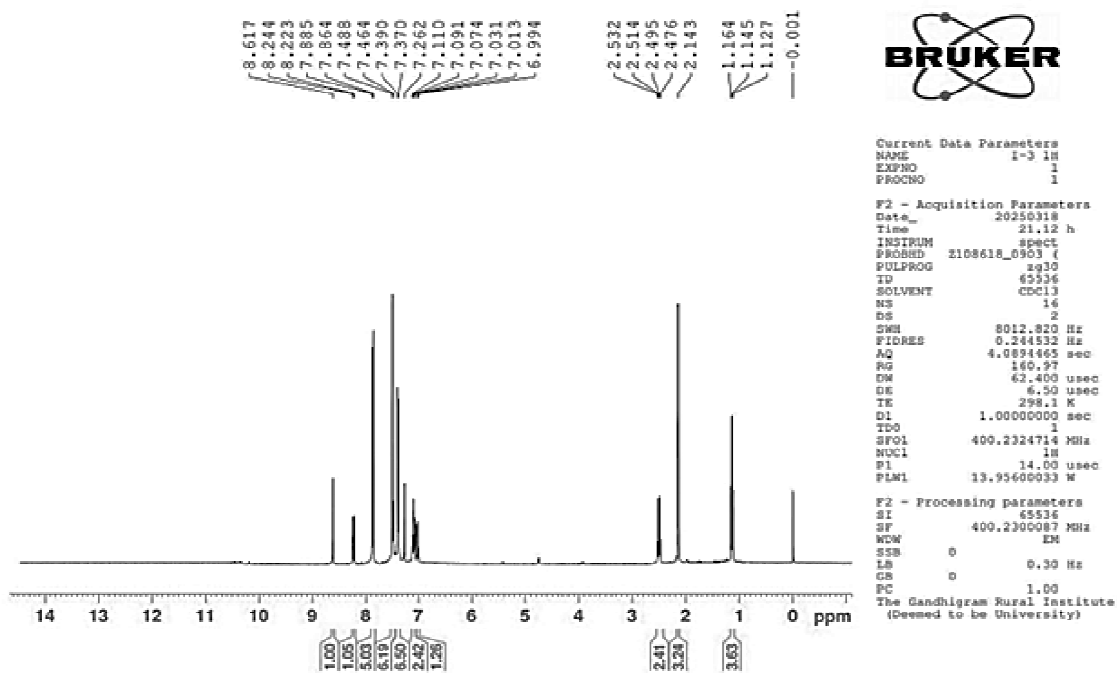


¹H NMR spectrum of (*E*)-*N*-(2-bromobenzylidene)-2-(trifluoromethyl)aniline (13)



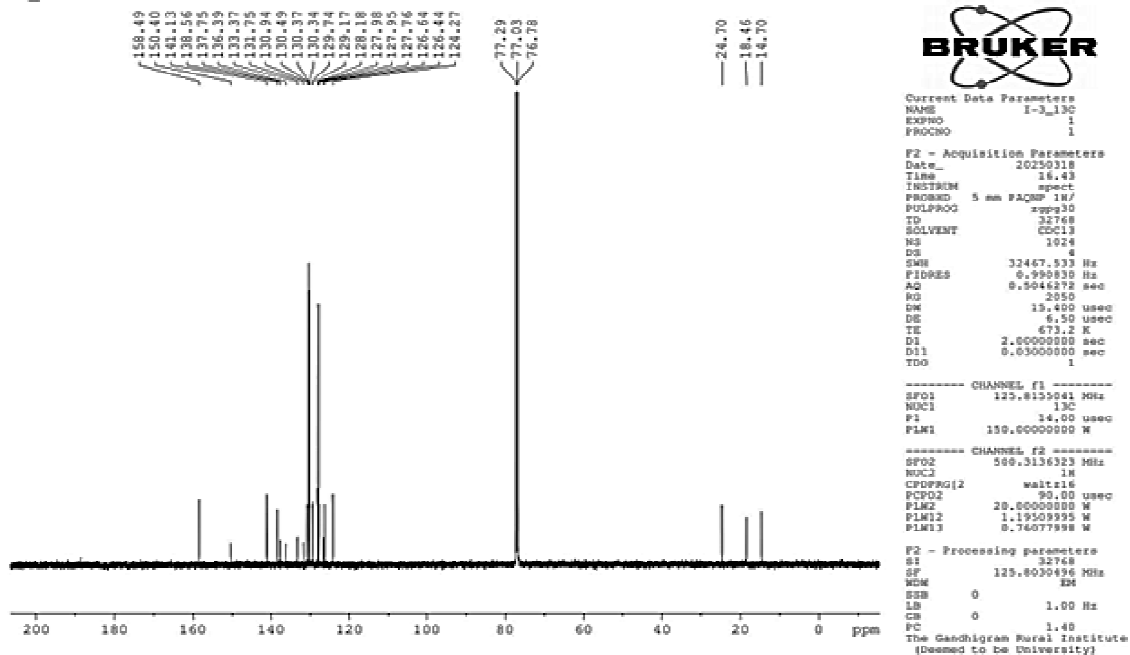
¹³C NMR spectrum of (*E*)-*N*-(2-bromobenzylidene)-2-(trifluoromethyl)aniline (13)

I-3 1H



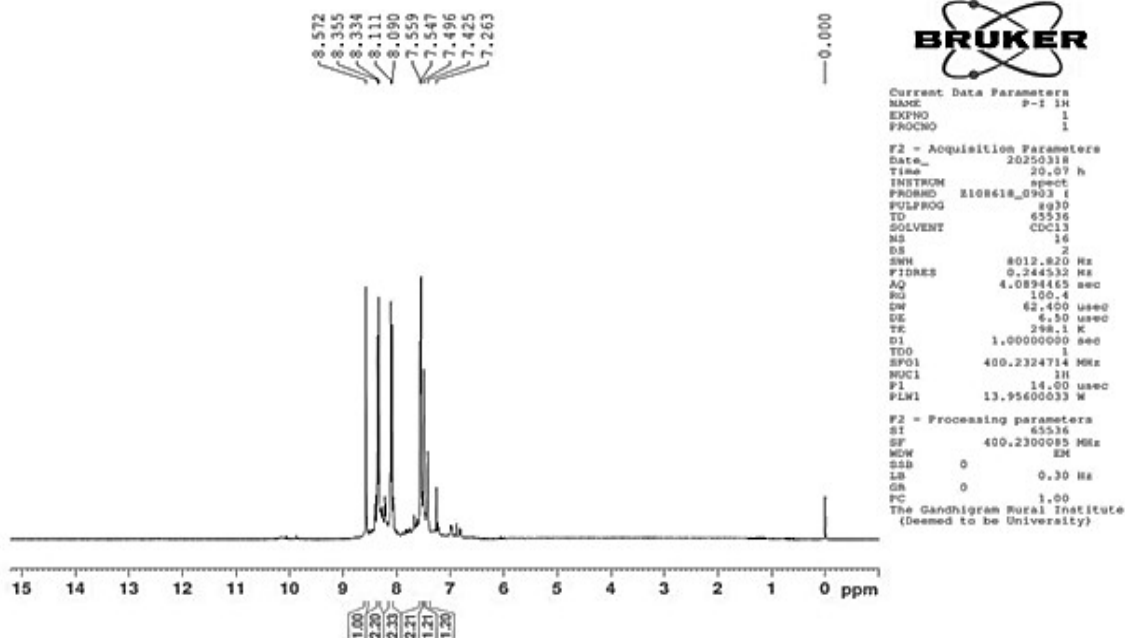
¹H NMR spectrum of (E)-N-(2,4-Dichlorobenzylidene)-2-ethyl-6-methylbenzenamine (15)

I-3_13C



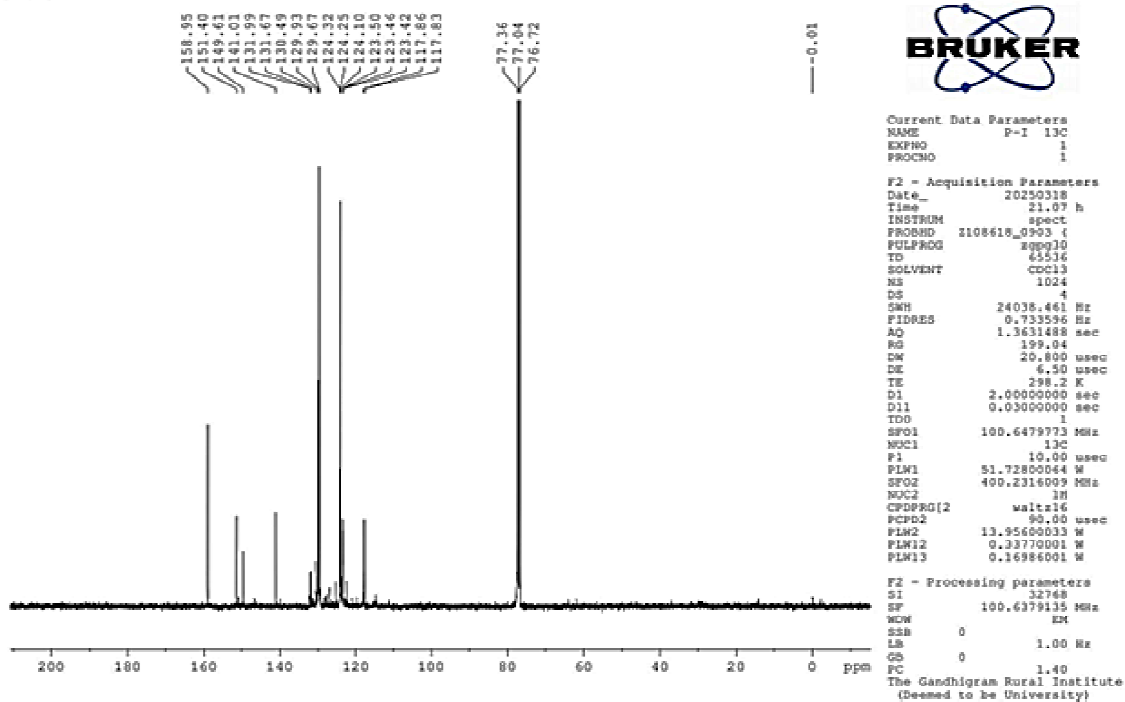
¹³C NMR spectrum of (E)-N-(2,4-Dichlorobenzylidene)-2-ethyl-6-methylbenzenamine (15)

P-I 1H



¹H NMR spectrum of (*E*)-*N*-(4-nitrobenzylidene)-3-(trifluoromethyl)benzenamine (16)

P-I 13C



¹³C NMR spectrum of (*E*)-*N*-(4-Nitrobenzylidene)-3-(trifluoromethyl)benzenamine (16)

checkCIF/PLATON report

Structure factors have been supplied for datablock(s) ATF2B_071124_0m_a

THIS REPORT IS FOR GUIDANCE ONLY. IF USED AS PART OF A REVIEW PROCEDURE FOR PUBLICATION, IT SHOULD NOT REPLACE THE EXPERTISE OF AN EXPERIENCED CRYSTALLOGRAPHIC REFEREE.

No syntax errors found. CIF dictionary Interpreting this report

Datablock: ATF2B_071124_0m_a

Bond precision: C-C = 0.0052 Å Wavelength=0.71073

Cell: a=10.0047(12) b=7.9137(10) c=16.739(2)
 alpha=90 beta=102.161(4) gamma=90

Temperature: 300 K

	Calculated	Reported
Volume	1295.6(3)	1295.6(3)
Space group	P 21/c	P 21/c
Hall group	-P 2ybc	-P 2ybc
Moiety formula	C14 H9 Br F3 N	C14 H9 Br F3 N
Sum formula	C14 H9 Br F3 N	C14 H9 Br F3 N
Mr	328.12	328.13
Dx, g cm ⁻³	1.682	1.682
Z	4	4
Mu (mm ⁻¹)	3.193	3.193
F000	648.0	648.0
F000'	647.23	
h, k, lmax	13, 10, 22	13, 10, 22
Nref	3224	3208
Tmin, Tmax	0.480, 0.629	0.552, 0.746
Tmin'	0.444	

Correction method= # Reported T Limits: Tmin=0.552 Tmax=0.746
AbsCorr = MULTI-SCAN

Data completeness= 0.995 Theta(max)= 28.300

R(reflections)= 0.0488(1888)

wR2(reflections)=
0.1285(3208)

S = 1.028

Npar= 172

The following ALERTS were generated. Each ALERT has the format

test-name_ALERT_alert-type_alert-level.

Click on the hyperlinks for more details of the test.



Alert level C

PLAT911_ALERT_3_C Missing FCF Refl Between Thmin & STh/L= 0.600 12 Report
3 0 0, -2 0 2, -2 1 2, -1 1 2, 1 0 2, 1 1 2,
2 0 2, 2 1 2, -2 0 4, 0 0 4, 1 0 4, 1 1 4,
PLAT913_ALERT_3_C Missing # of Very Strong Reflections in FCF 6 Note
3 0 0, 2 1 2, -2 0 4, 0 0 4, 1 0 4, 1 1 4,



Alert level G

PLAT242_ALERT_2_G Low 'MainMol' Ueq as Compared to Neighbors of C14 Check
PLAT883_ALERT_1_G No Info/Value for _atom_sites_solution_primary . Please Do !
PLAT912_ALERT_4_G Missing # of FCF Reflections Above STh/L= 0.600 4 Note
PLAT969_ALERT_5_G The 'Henn et al.' R-Factor-gap value 3.755 Note
Predicted wR2: Based on SigI**2 3.42 or SHELX Weight 12.50
PLAT978_ALERT_2_G Number C-C Bonds with Positive Residual Density. 3 Info

- 0 **ALERT level A** = Most likely a serious problem - resolve or explain
0 **ALERT level B** = A potentially serious problem, consider carefully
2 **ALERT level C** = Check. Ensure it is not caused by an omission or oversight
5 **ALERT level G** = General information/check it is not something unexpected
- 1 ALERT type 1 CIF construction/syntax error, inconsistent or missing data
2 ALERT type 2 Indicator that the structure model may be wrong or deficient
2 ALERT type 3 Indicator that the structure quality may be low
1 ALERT type 4 Improvement, methodology, query or suggestion
1 ALERT type 5 Informative message, check
-
-

It is advisable to attempt to resolve as many as possible of the alerts in all categories. Often the minor alerts point to easily fixed oversights, errors and omissions in your CIF or refinement strategy, so attention to these fine details can be worthwhile. In order to resolve some of the more serious problems it may be necessary to carry out additional measurements or structure refinements. However, the purpose of your study may justify the reported deviations and the more serious of these should normally be commented upon in the discussion or experimental section of a paper or in the "special_details" fields of the CIF. checkCIF was carefully designed to identify outliers and unusual parameters, but every test has its limitations and alerts that are not important in a particular case may appear. Conversely, the absence of alerts does not guarantee there are no aspects of the results needing attention. It is up to the individual to critically assess their own results and, if necessary, seek expert advice.

Publication of your CIF in IUCr journals

A basic structural check has been run on your CIF. These basic checks will be run on all CIFs submitted for publication in IUCr journals (*Acta Crystallographica*, *Journal of Applied Crystallography*, *Journal of Synchrotron Radiation*); however, if you intend to submit to *Acta Crystallographica Section C* or *E* or *IUCrData*, you should make sure that full publication checks are run on the final version of your CIF prior to submission.

Publication of your CIF in other journals

Please refer to the *Notes for Authors* of the relevant journal for any special instructions relating to CIF submission.

