

## Supplementary Materials

Figure S1(a)

Full Data set

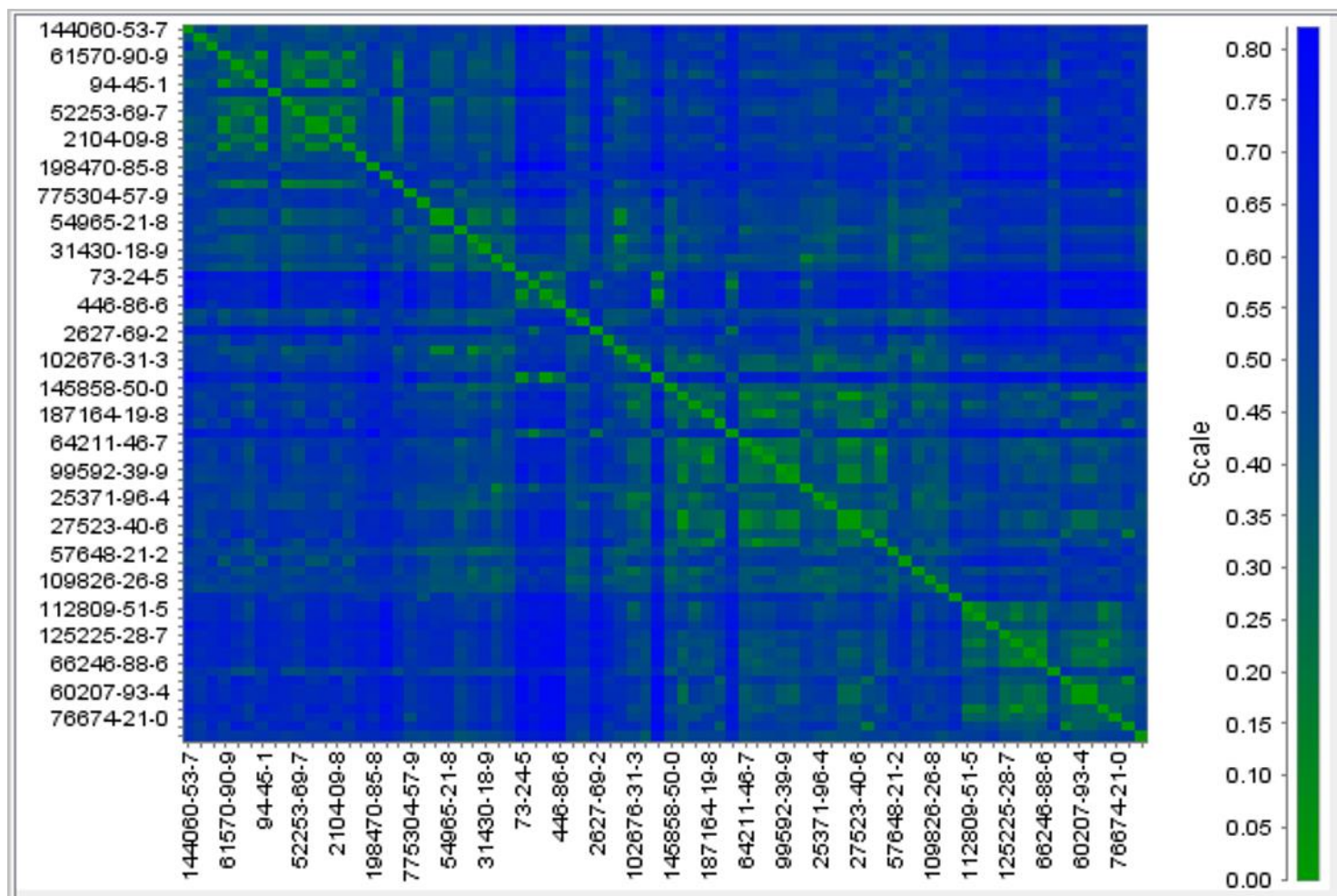
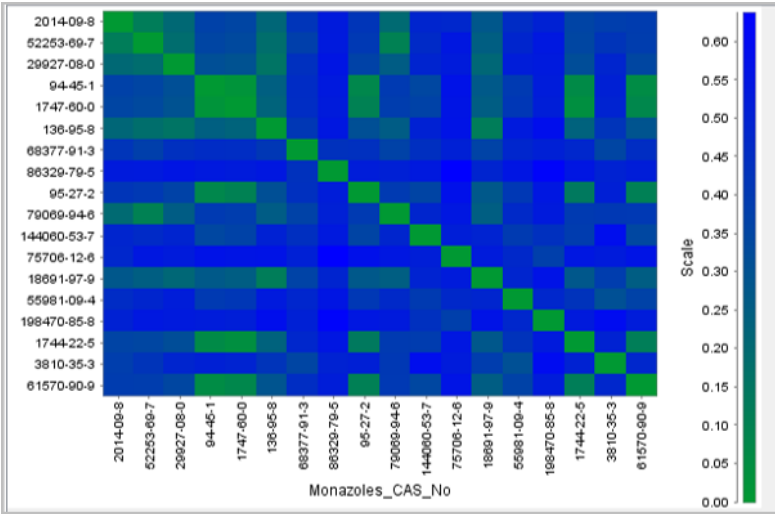
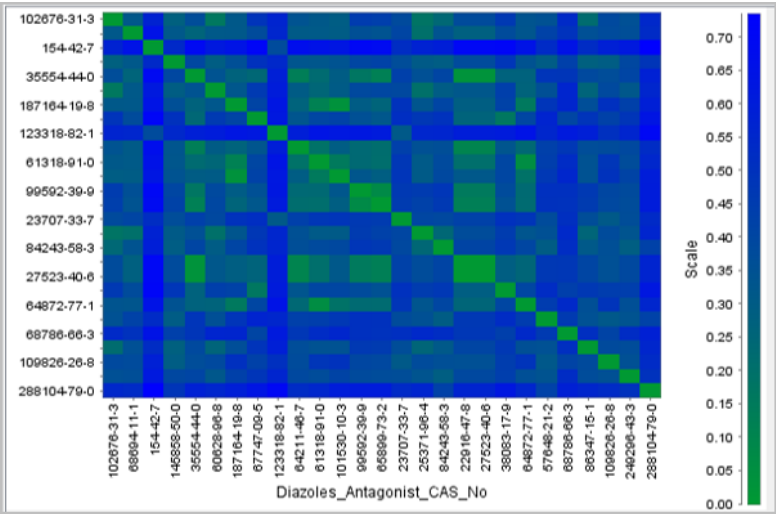


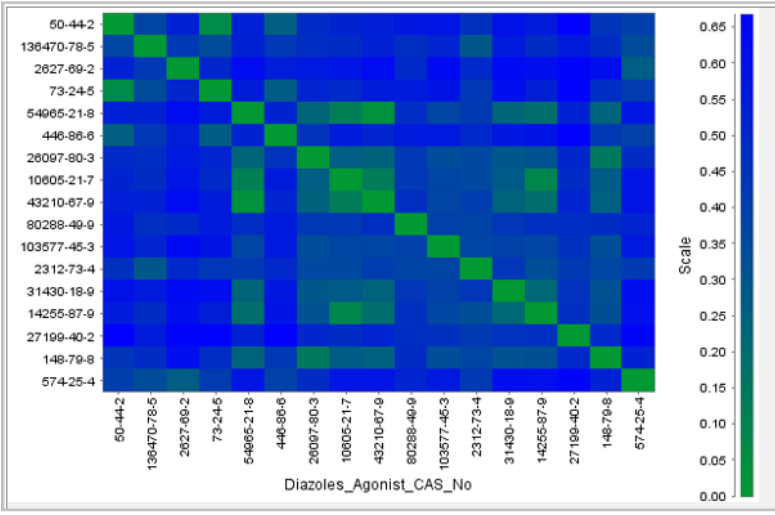
Figure S1(b)



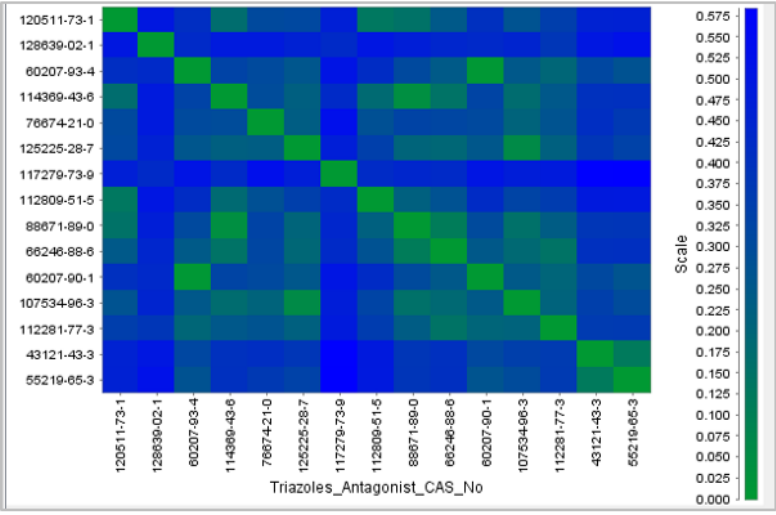
(c)



(d)



(e)

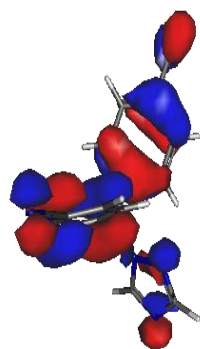


(f)

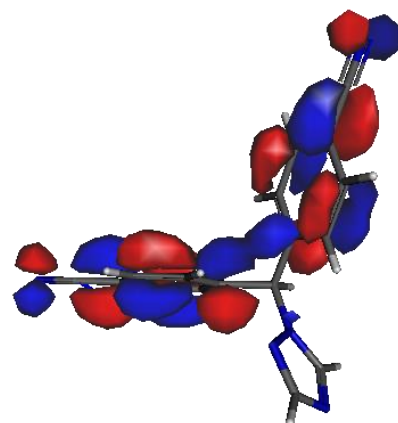
**Figure S1:** The heat maps showing the diversity of the data set. a.) Heat map for full data set. b) Heat map for the classified data set based on activity and class of the azole. On left side agonist monazoles (Thiazole/oxazole) (c) and agonist diazoles (imidazoles and benzimidazole) (e). On right side antagonist diazoles (imidazoles and benzimidazole) (d) and antagonist triazoles (f).

Molecule  
Litrazole

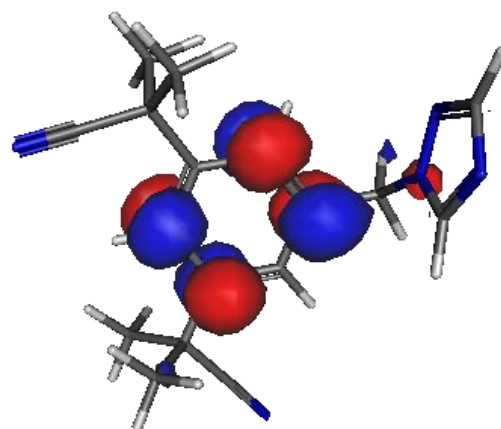
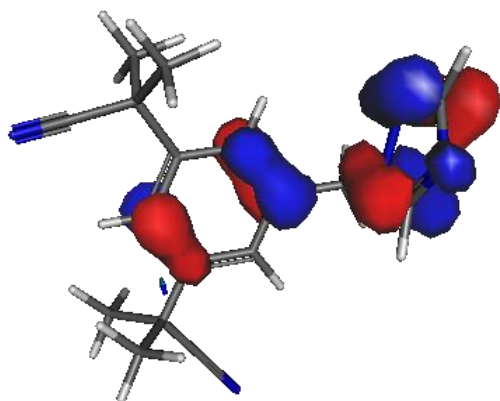
HOMO



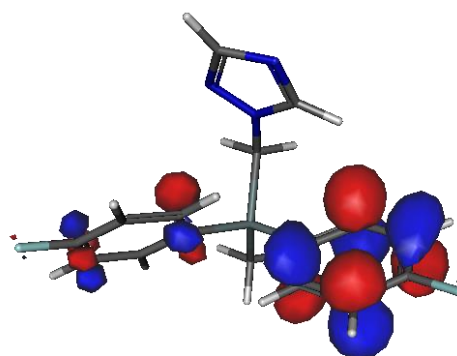
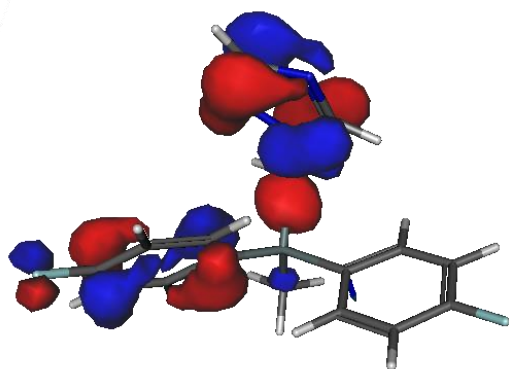
LUMO



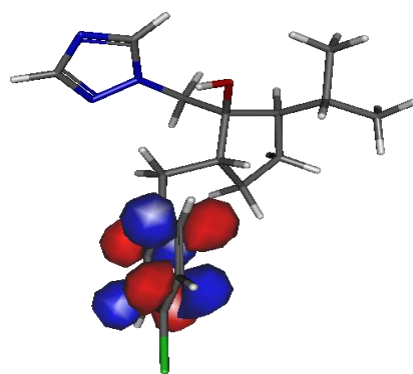
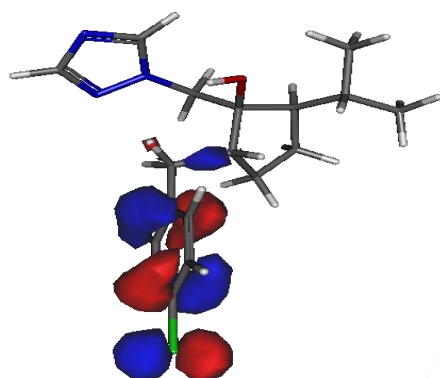
Anastrozole



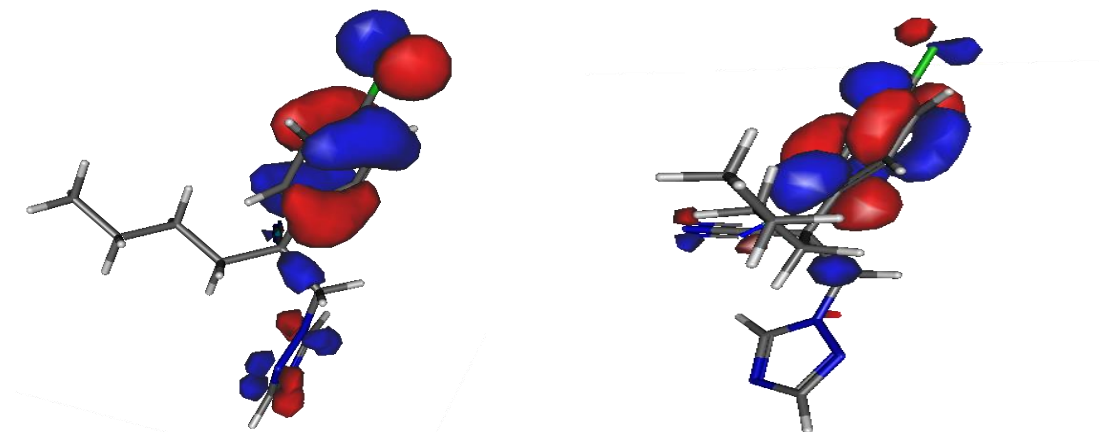
Flusilazole



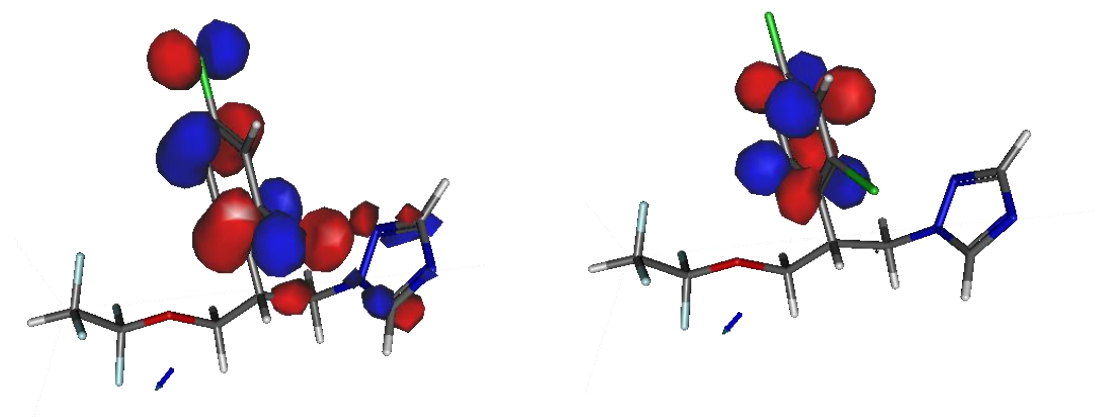
Ipiconazole



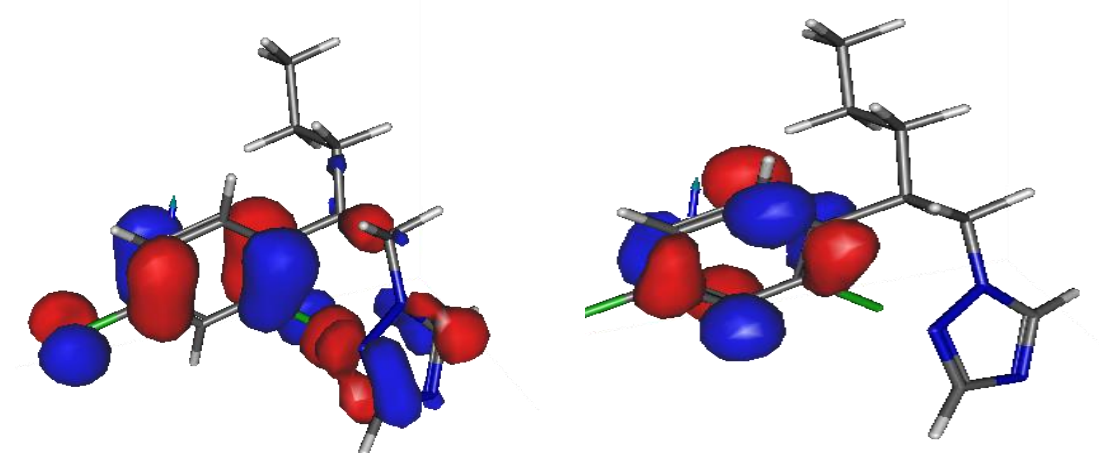
Myclobutanil



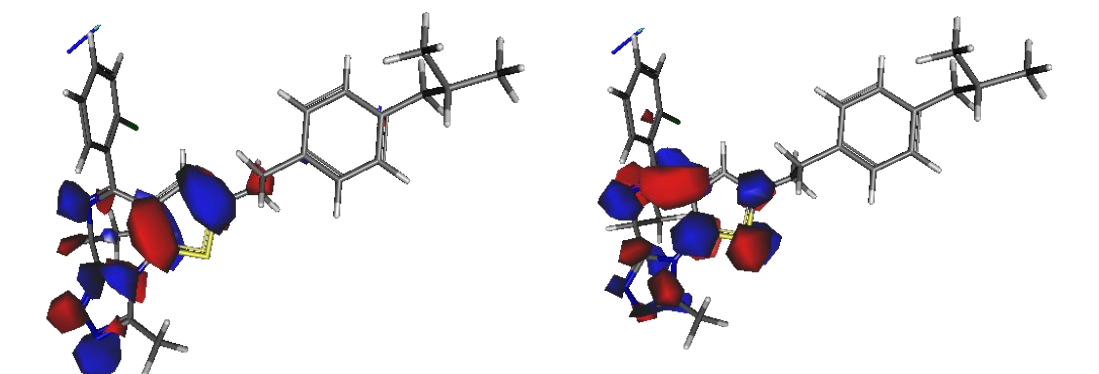
Tetraconazole



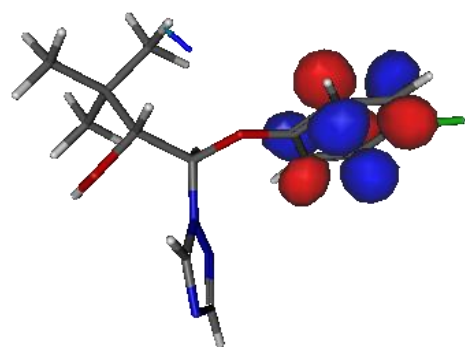
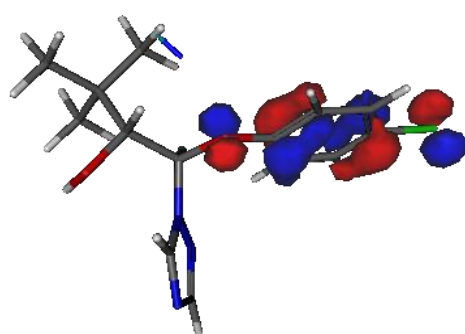
Penconazole



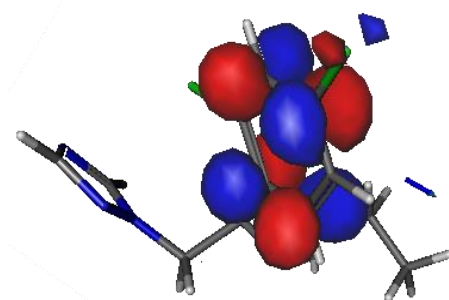
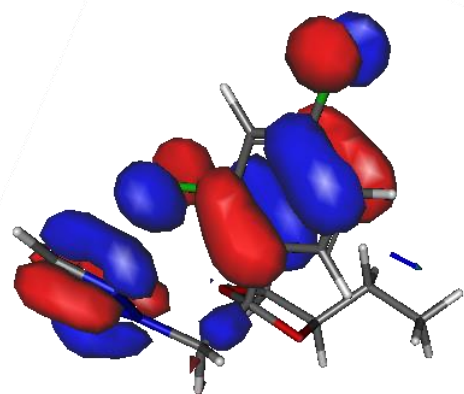
Israpafant



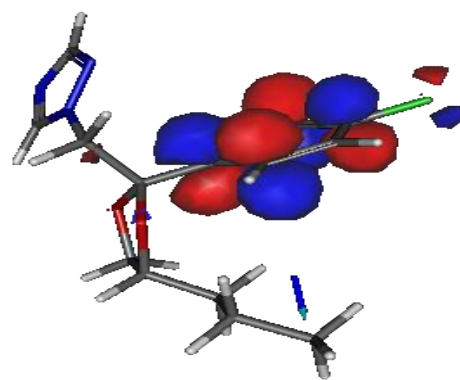
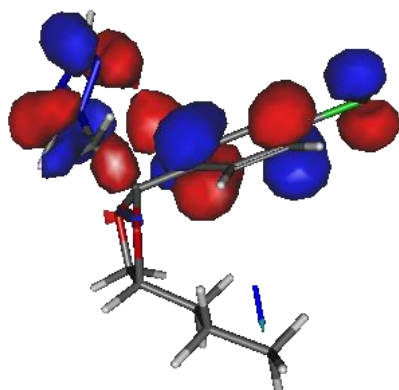
Triadimenol



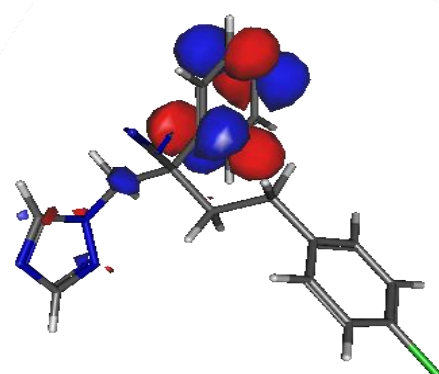
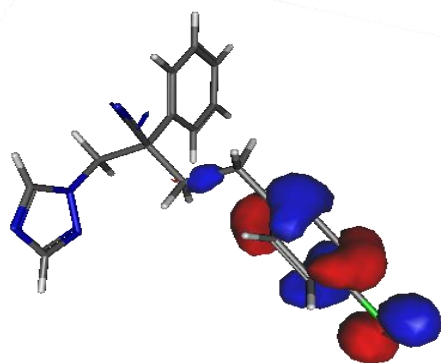
Etaconazole



Propiconazole

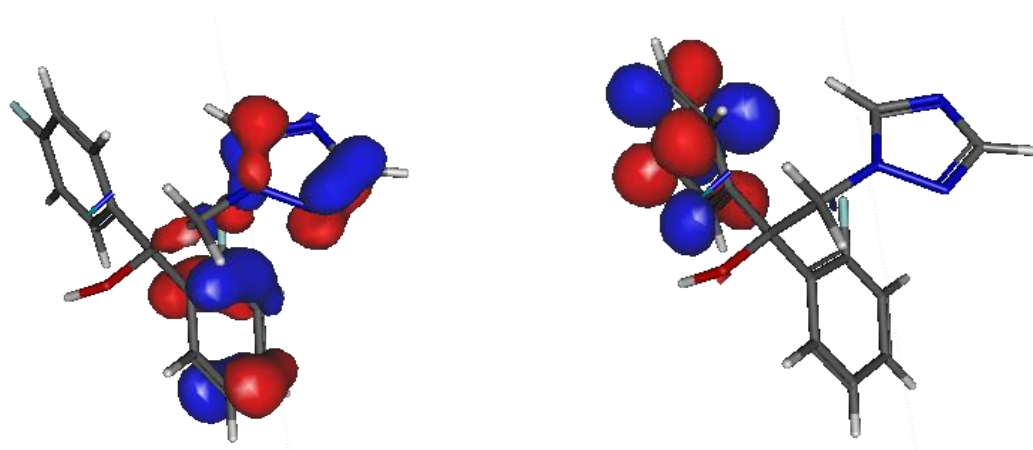


Fenbuconazole

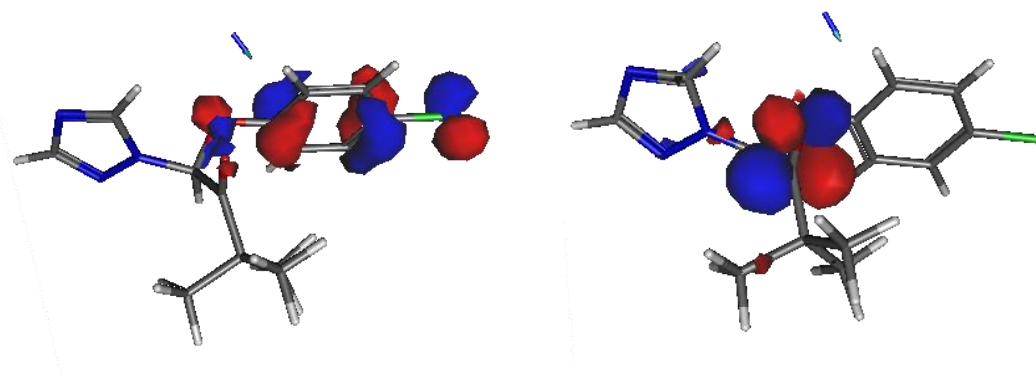




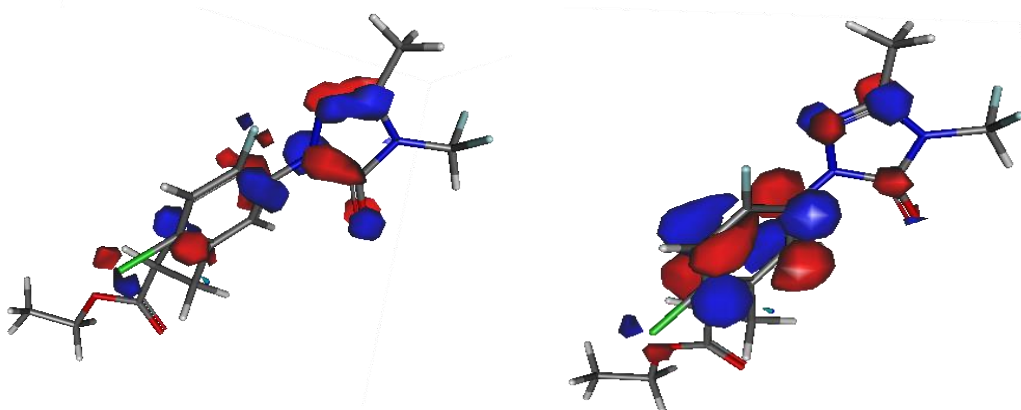
Flutriafol



Triadimefon



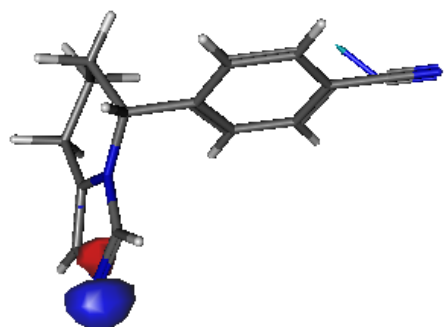
Carfentrazone-ethyl



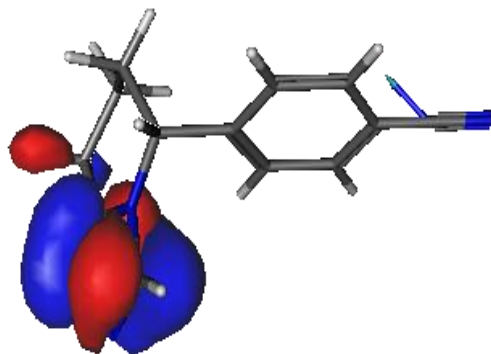
**Figure S2:** Representations of HOMO and LUMO isosurfaces for the antagonist triazoles computed using the single point at B3LYP/def2-SV(P) method on the structure obtained following complete optimisation with PM7.

Fadrozole

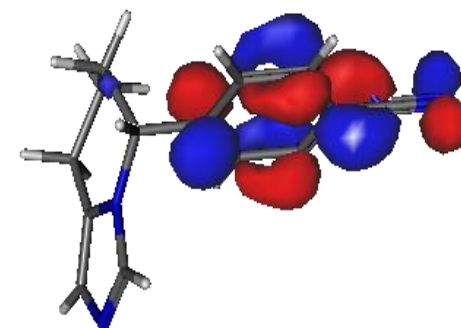
HOMONL



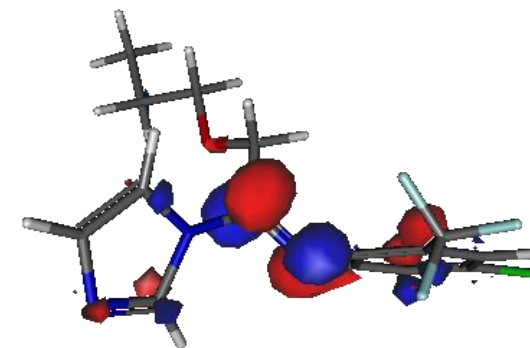
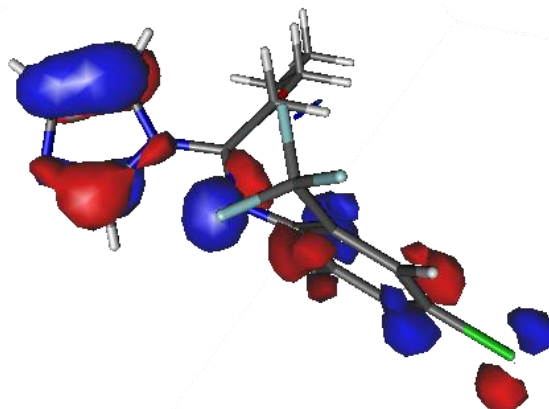
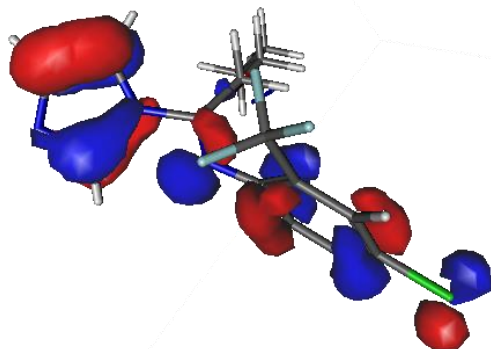
HOMO



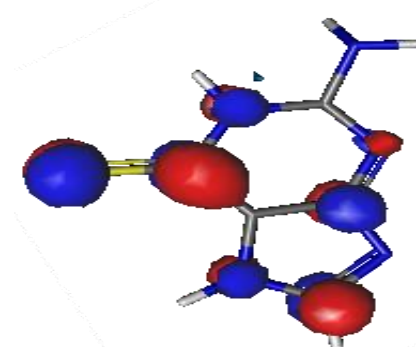
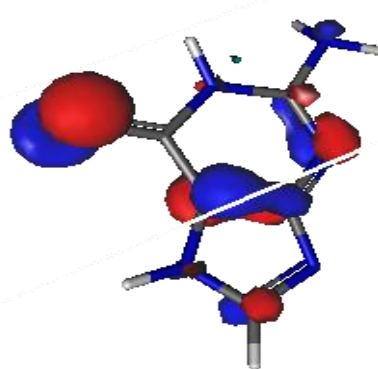
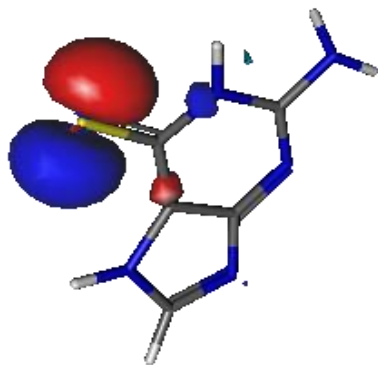
LUMO



Triflumizole

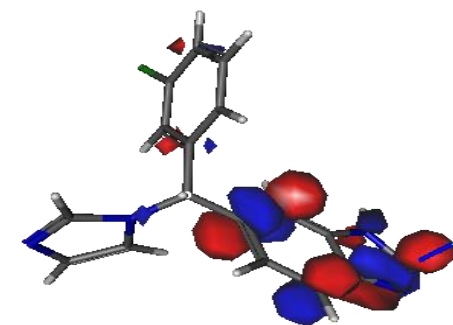
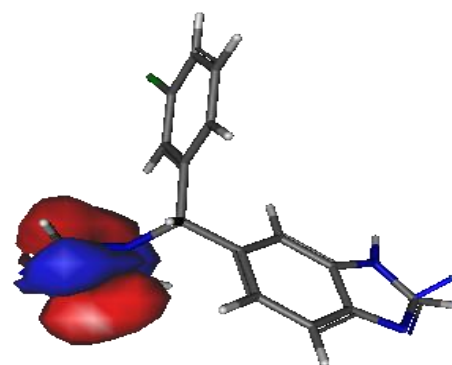
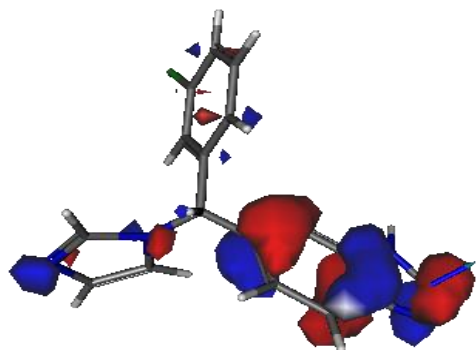


6-Thioguanine

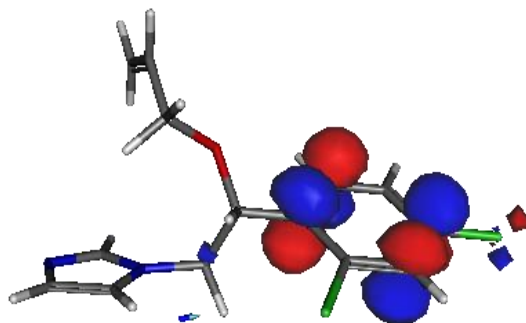
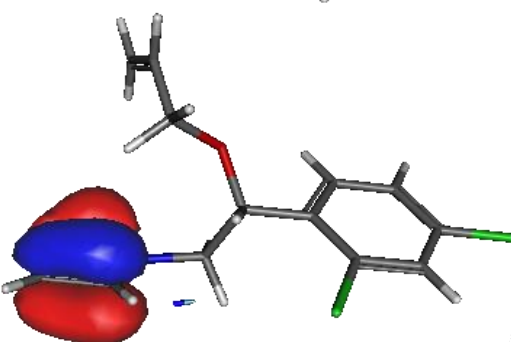
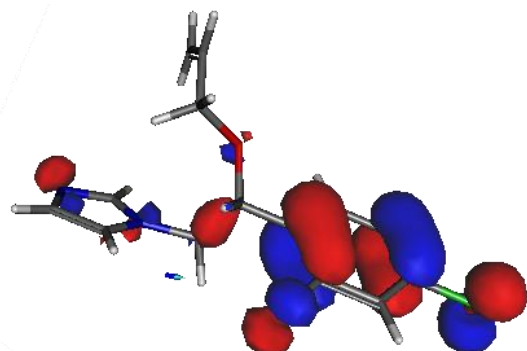




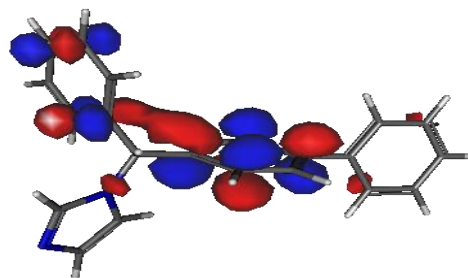
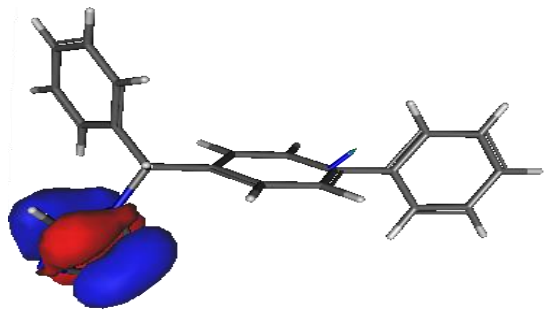
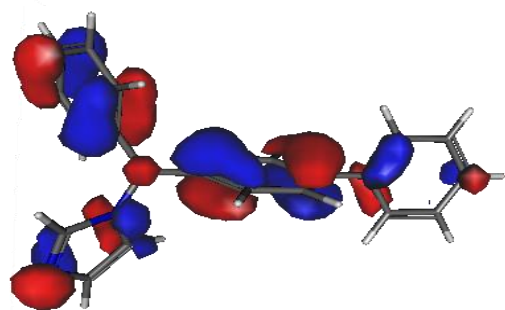
Liarozole



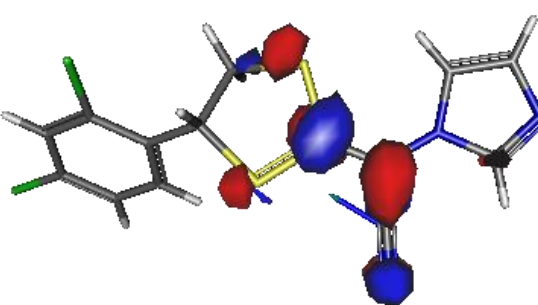
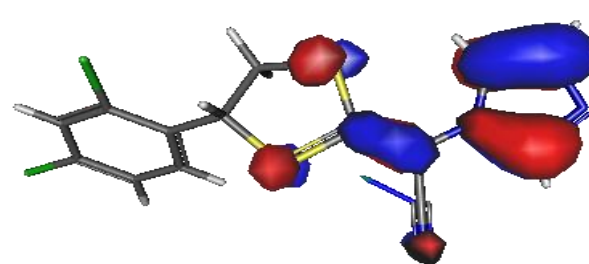
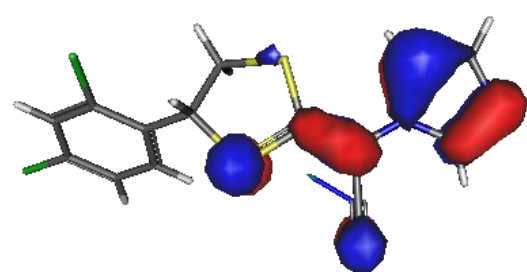
Imazalil



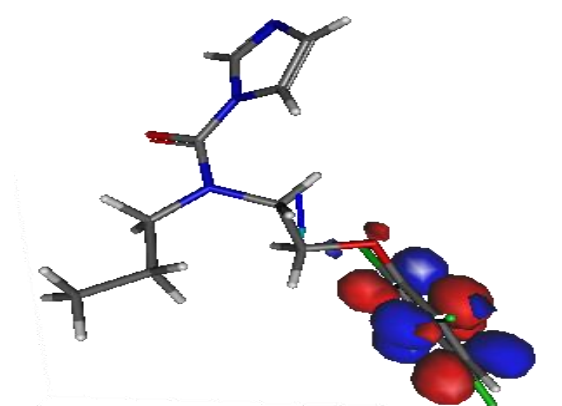
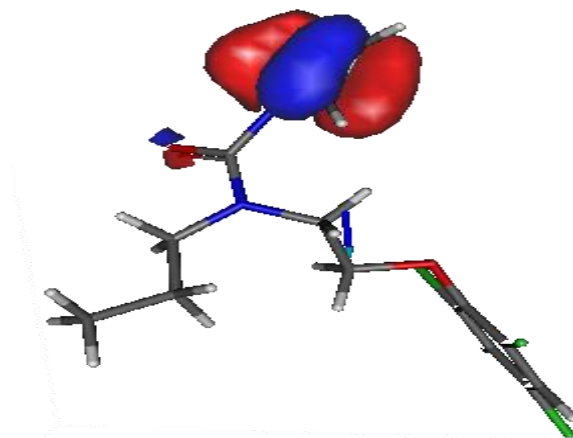
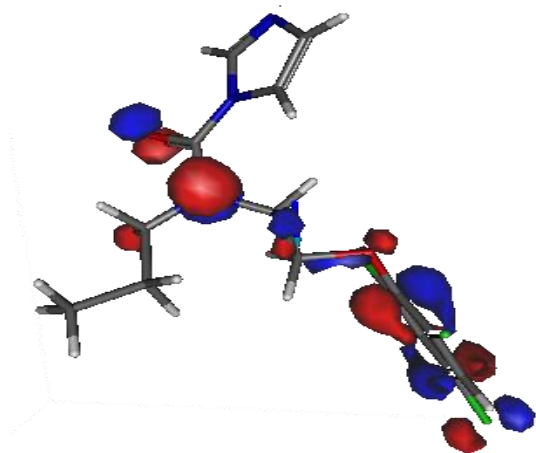
Bifonazole



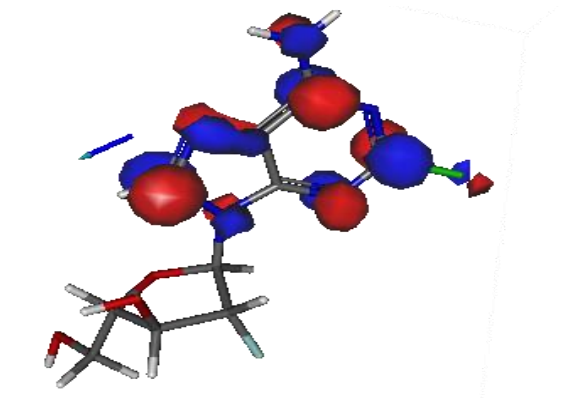
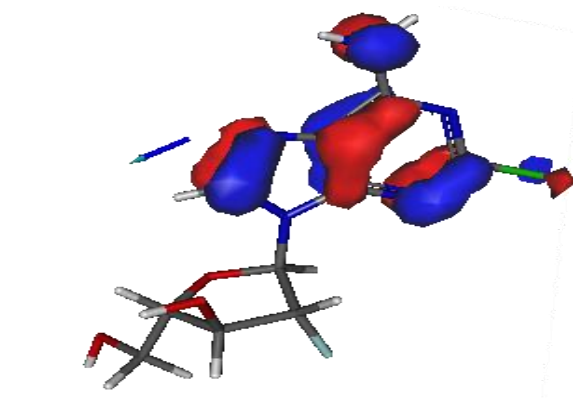
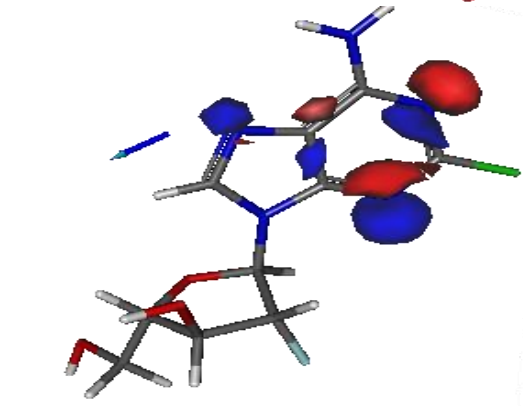
Luliconazole



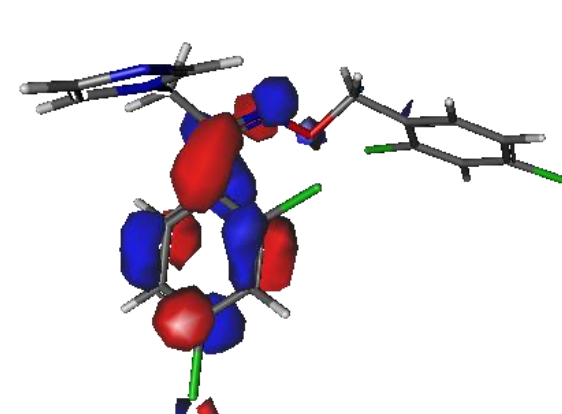
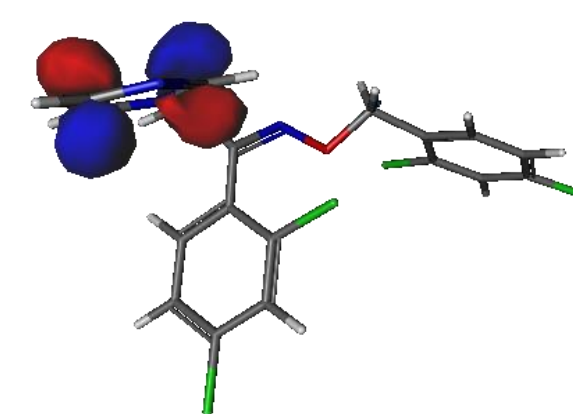
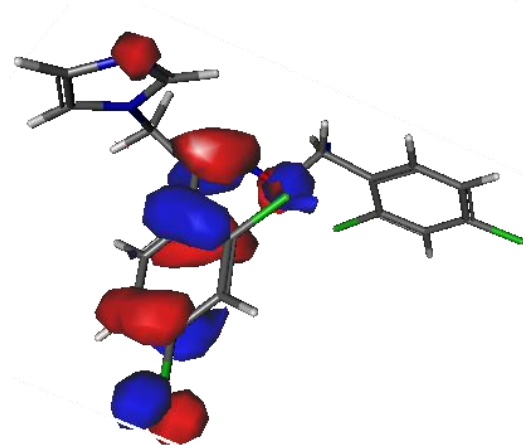
Prochloraz



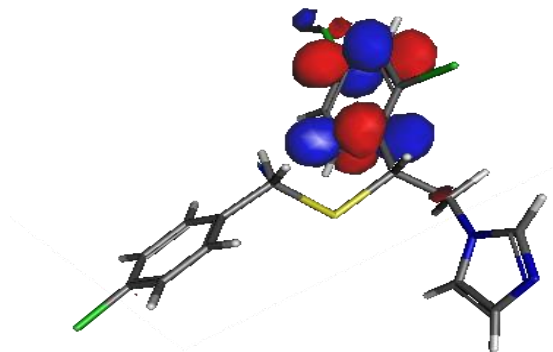
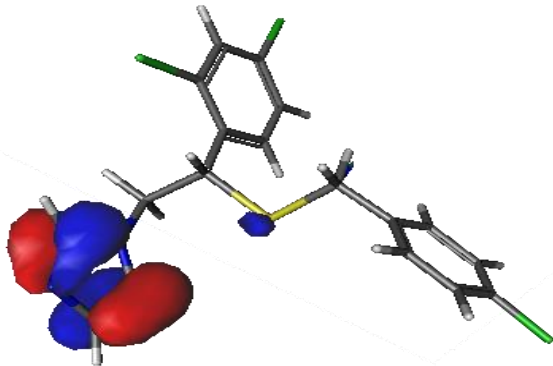
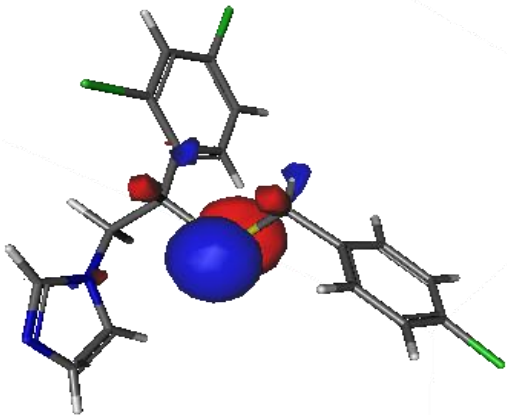
Clofarabine



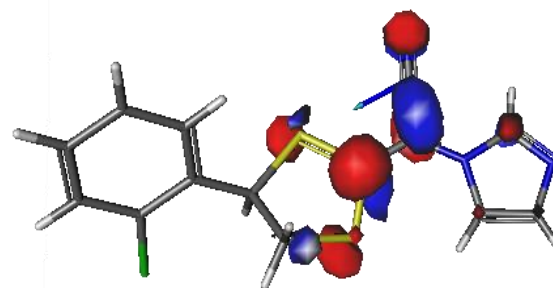
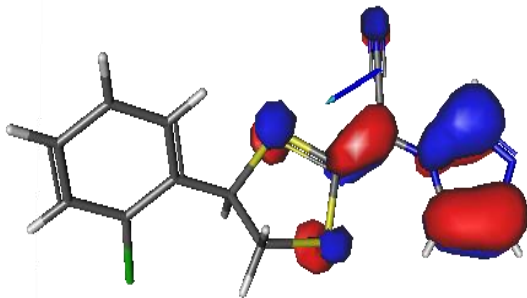
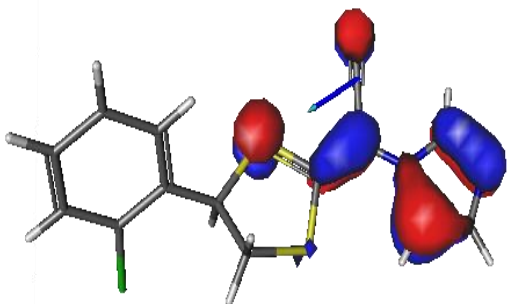
Oxiconazole



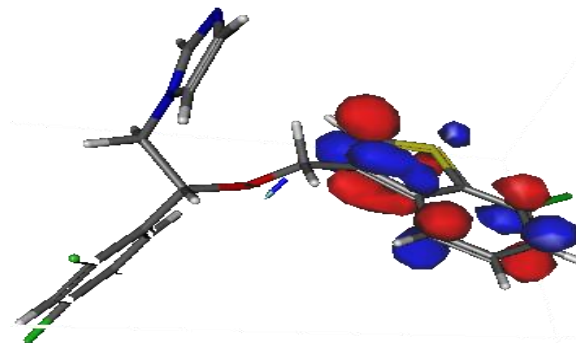
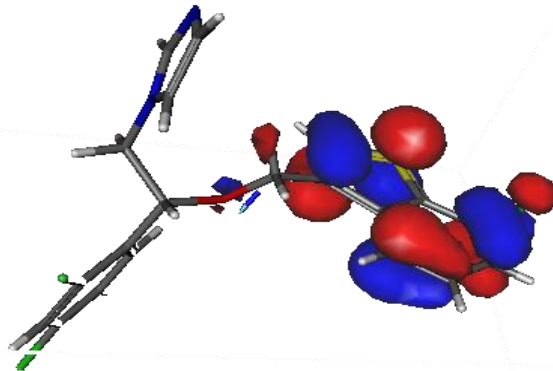
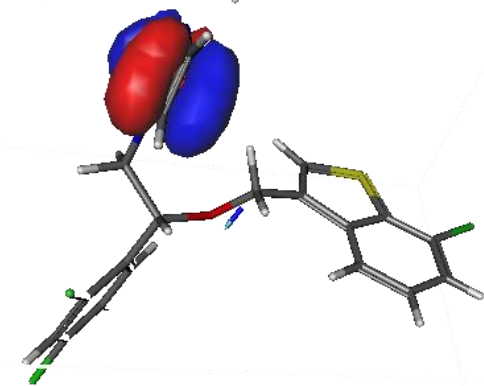
Sulconazole



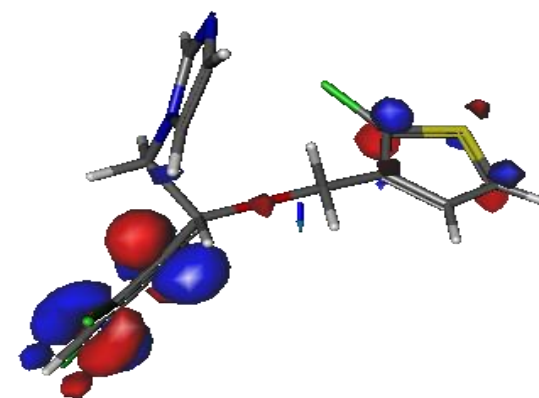
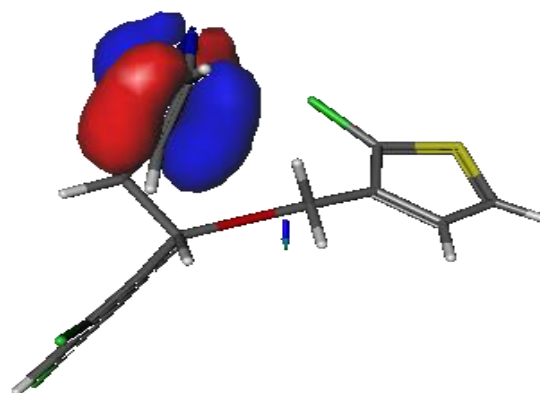
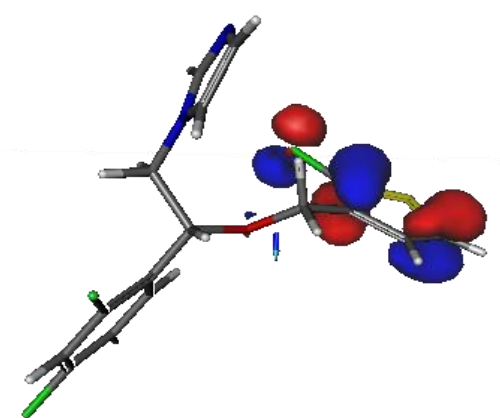
Lanoconazole



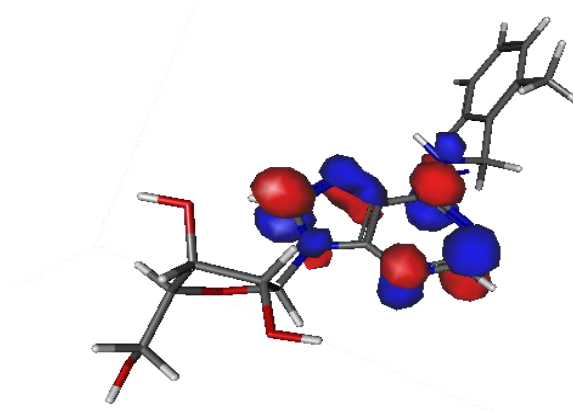
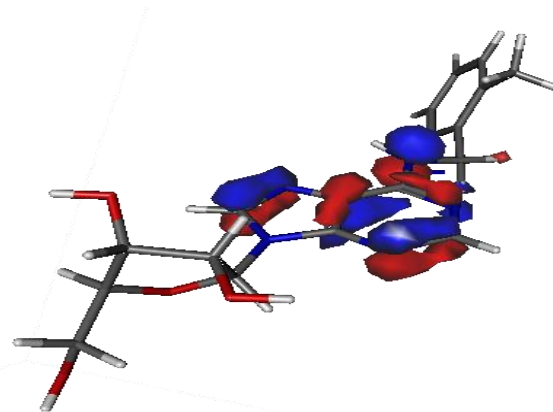
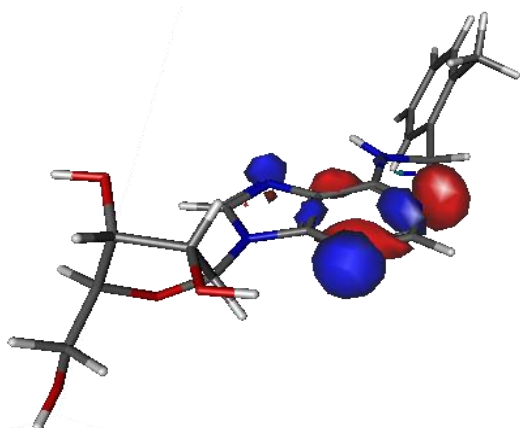
Sertaconazol



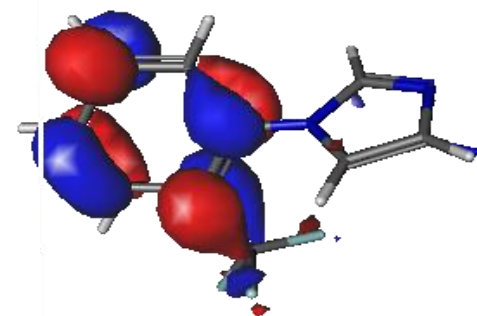
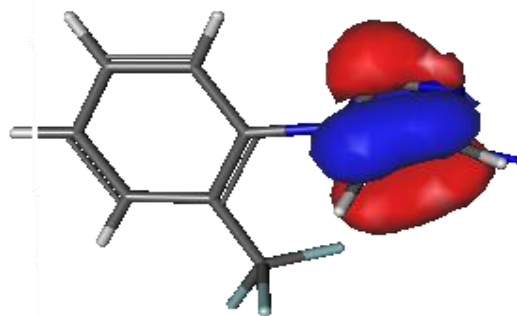
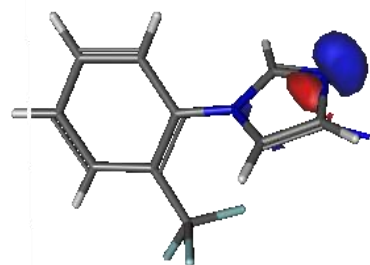
Tioconazole



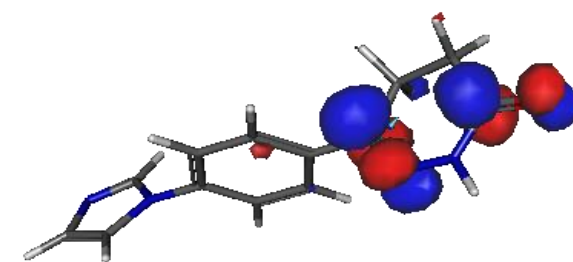
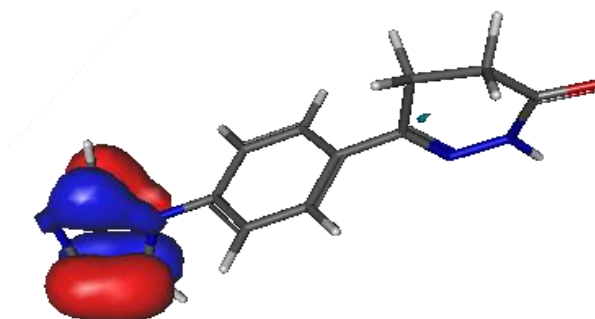
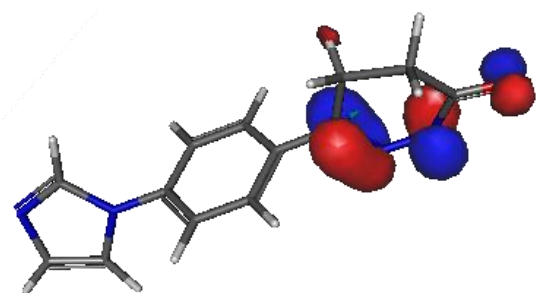
Metrifudil



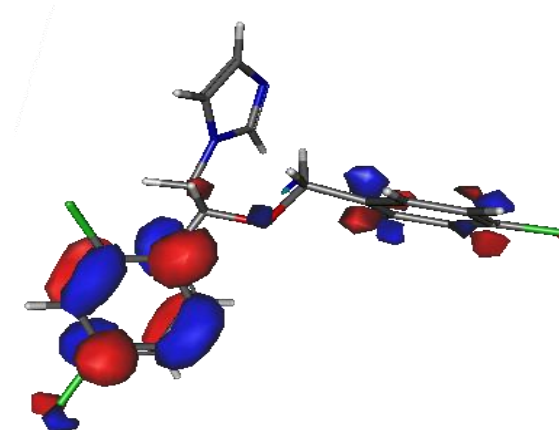
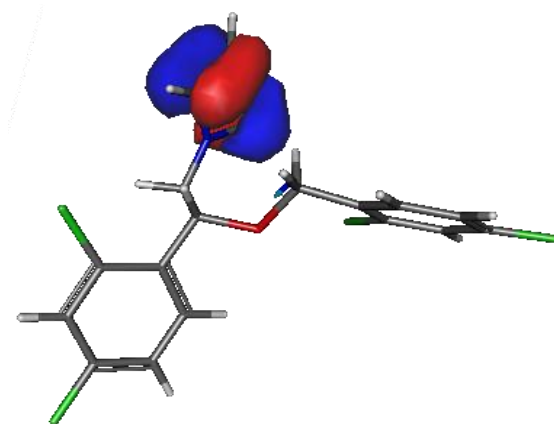
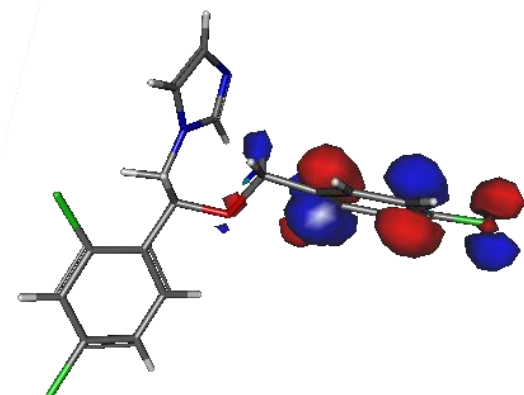
1-[2-(Trifluoromethyl)phenyl]-1H-imidazole



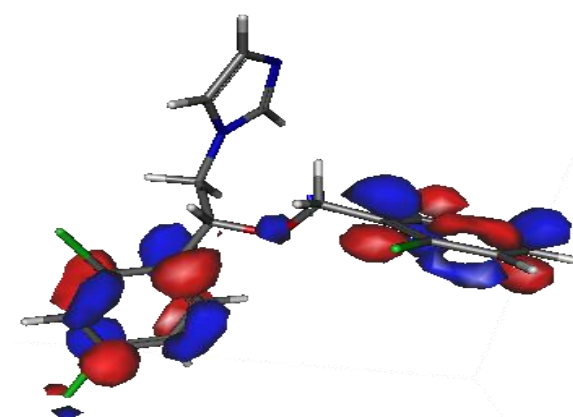
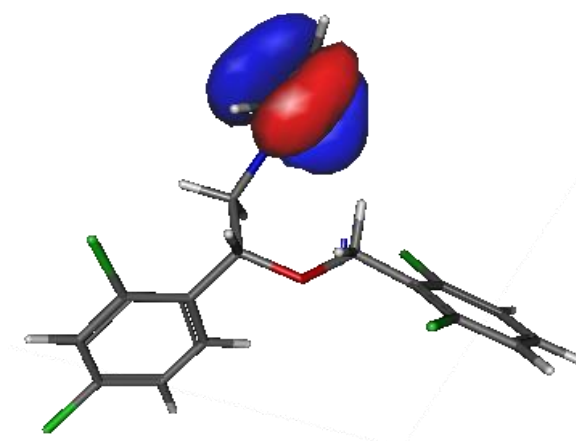
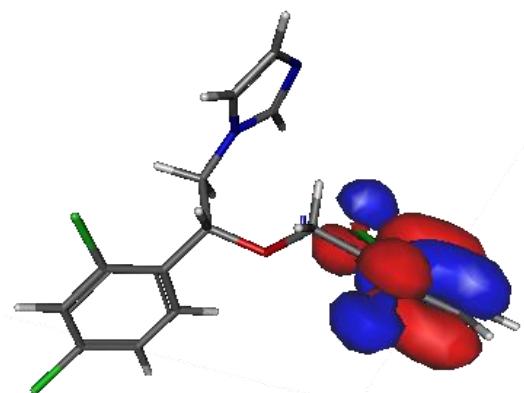
Imazodan



Miconazole

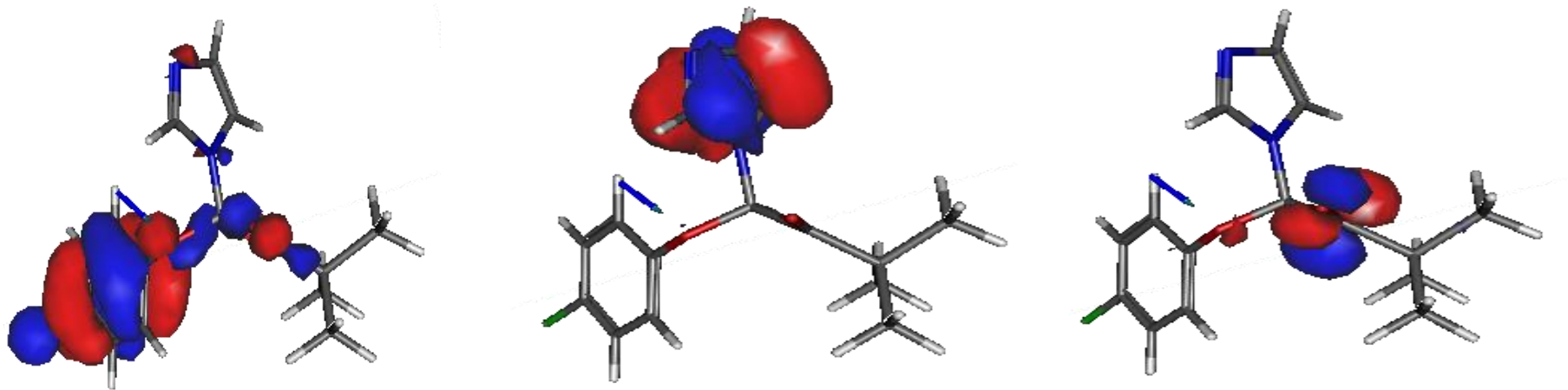


Isoconazole

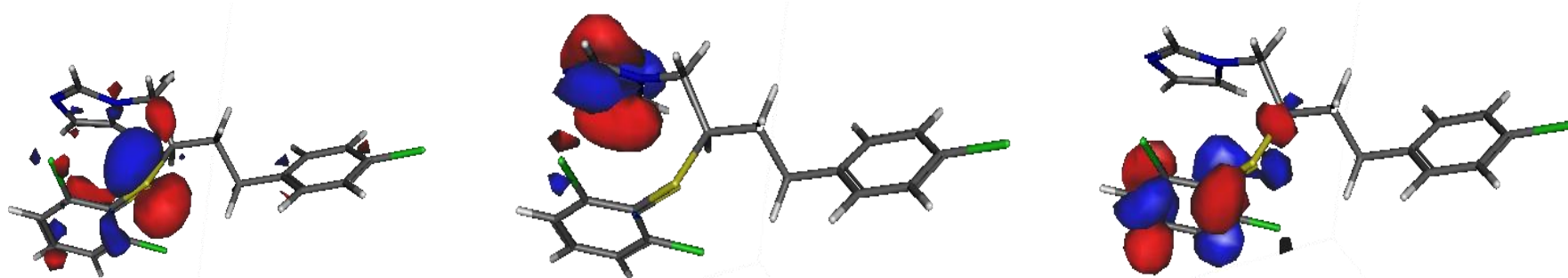




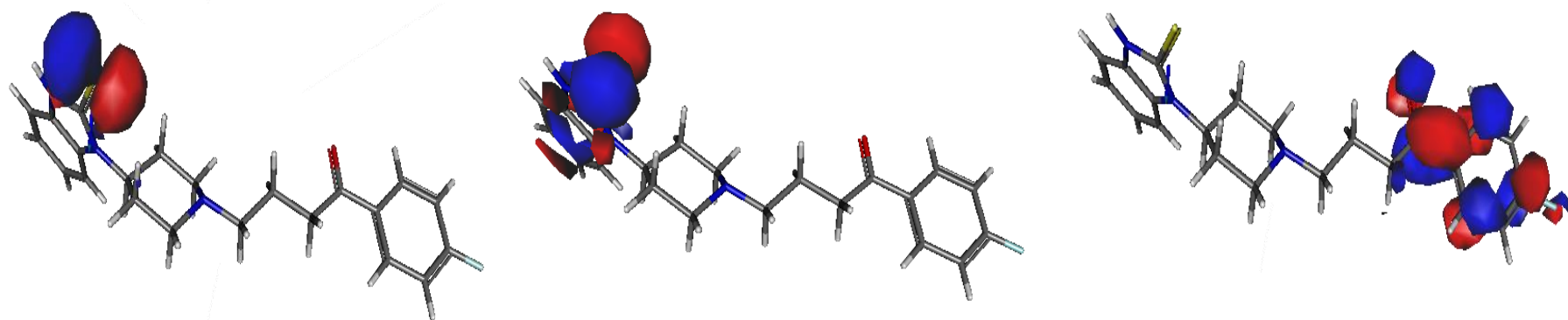
Climbazole



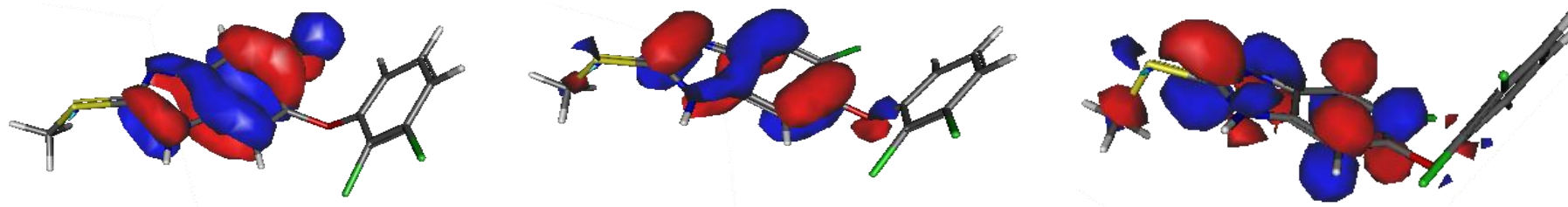
Butoconazole



Timiperone

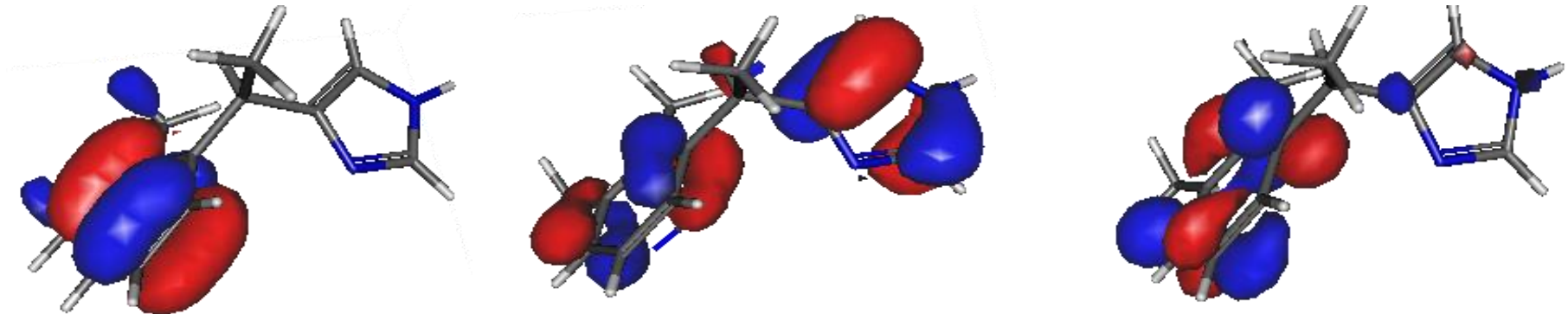


Triclabendazole

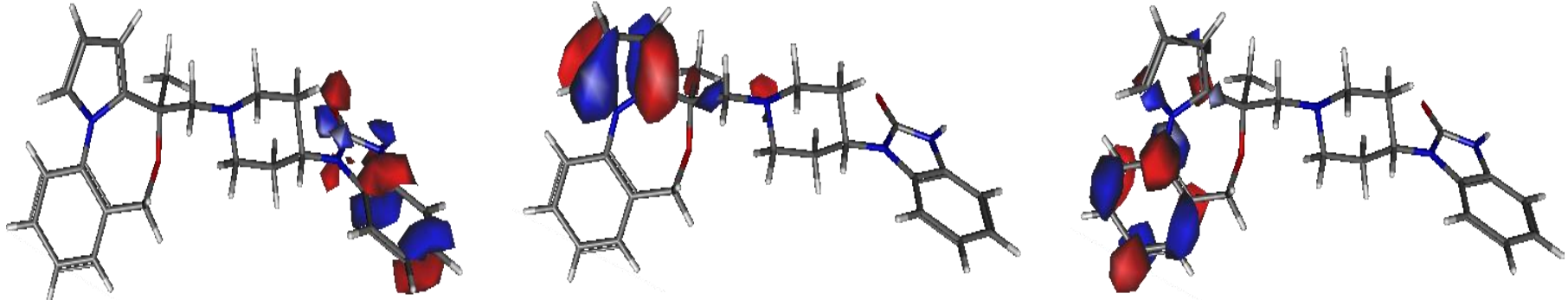




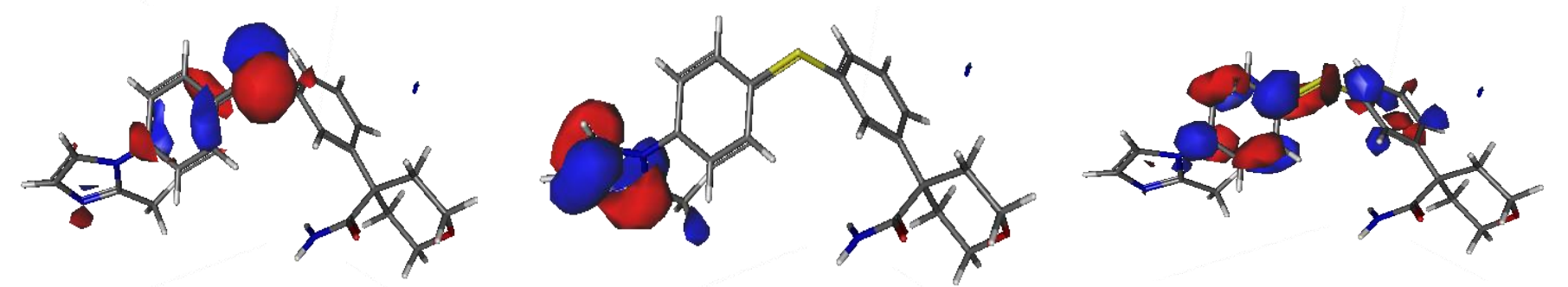
Medetomidine



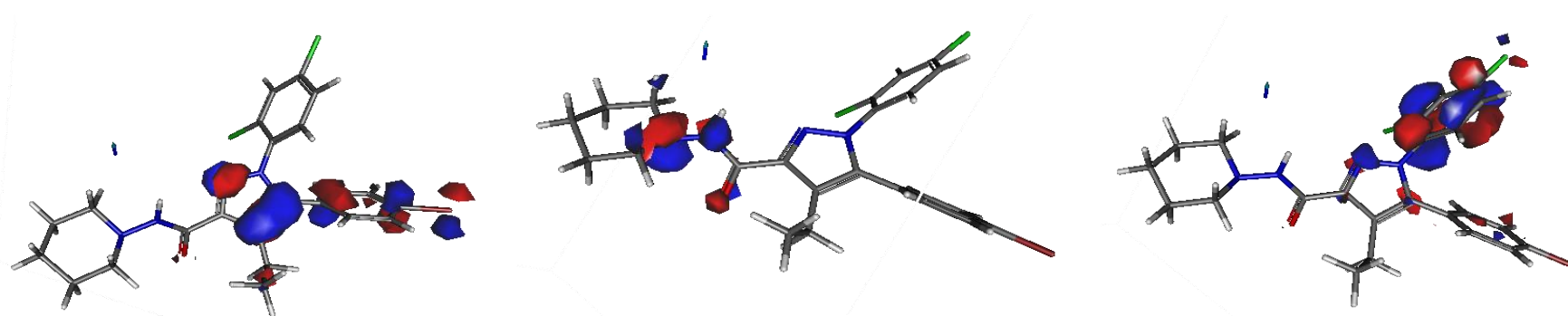
Zaldaride



CJ-013610

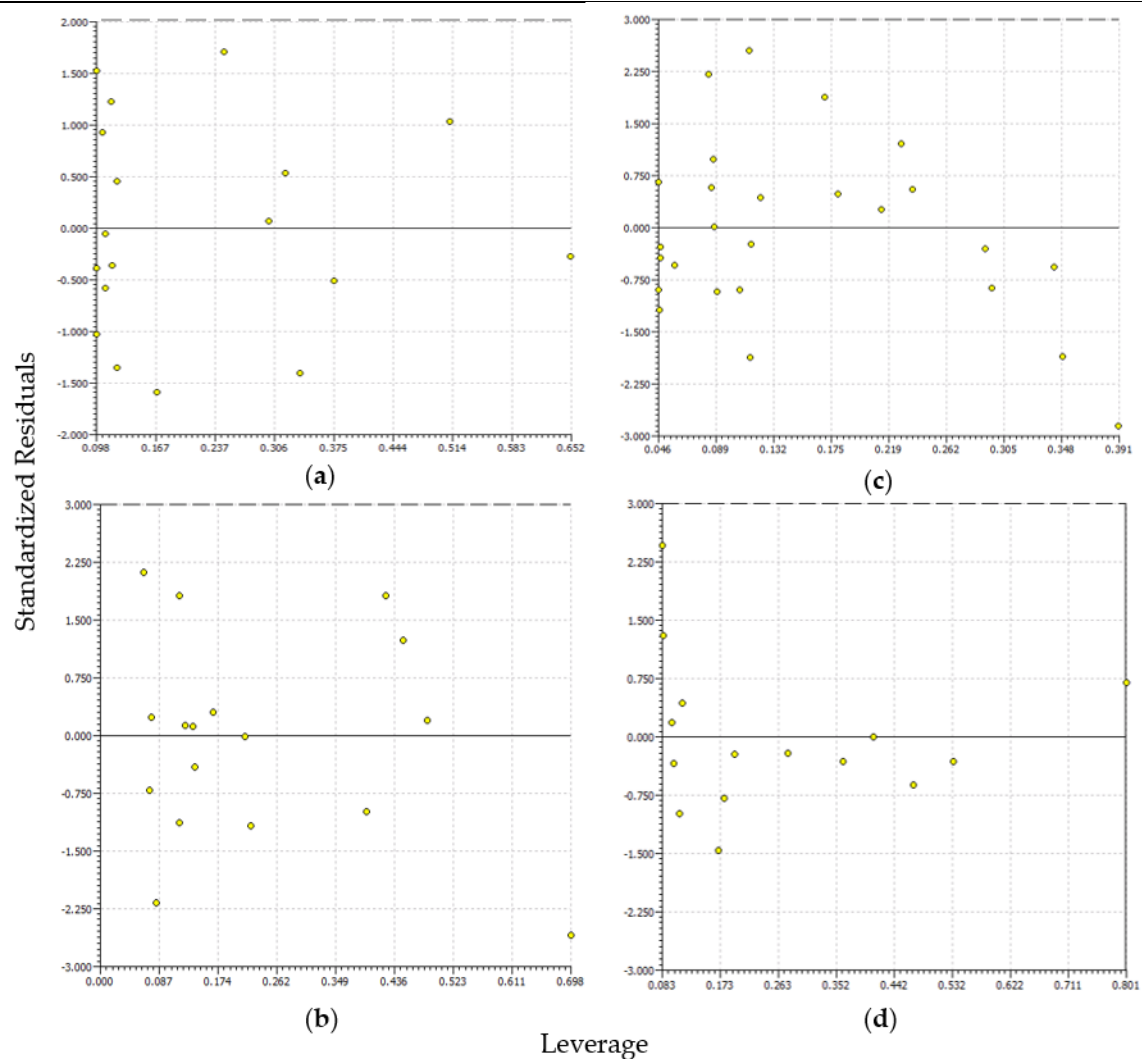


Surinabant



**Figure S3:** Representation of HOMO-1 (HOMONL), HOMO and LUMO isosurfaces of antagonist diazoles computed using the single point at B3LYP/def2-TZVP method on the structure obtained following complete optimisation with PM7

**Figure S4**



**Figure S4.** Williams plots of leverage vs standardised residuals for equations (1, 2, 4, 5) for: (a) agonist monazoles (*Thiazole/oxazole*); (b) agonist Diazoles (*imidazoles* and *benzimidazole*); (c) antagonist Diazoles (*imidazoles* and *benzimidazole*); (d) antagonist Triazoles.