



International Journal of ChemTech Research CODEN (USA): IJCRGG, ISSN: 0974-4290, ISSN(Online):2455-9555 Vol.12 No.04, pp 277-282, 2019

Different classical hydrogen –bonding patterns in two imidazole derivatives (N-(tert-butyl)-2-(4nitrophenyl)imidazo[1,2-a]pyrazin-3-amine and N-(tertbutyl)-2-(2-nitrophenyl)imidazo[1,2-a]pyridin-3-amine)

K. Hemanathan¹, R. Raja², K. Sakthi Murugesan^{1*}

^{1, 2, 1*}Department of Physics, Presidency College (Autonomous), Chennai - 600 005, India

Abstract : The title compounds, $C_{16}H_{17}N_5O_2$ in (I) and $C_{17}H_{18}N_4O_4$ in (II), are imidazole derivatives in which the imidazole moiety is fused with a piperdine ring system in (I), and with an benzene ring system in (II). The compound I, crystallized in the triclinic space group p1-with two molecules in a unit cell and compound II, crystallizes in a monoclinic in a C-centered lattice with eight molecules in the unit cell. According to single-crystal x-ray data, intra (C-H...N 3.397(3)) Å and intermolecular hydrogen bonds C-H...O 3.397(3)Å are formed between C-H groups of the imidazole cycle and O atoms of the paramagnetic moieties. The intermolecular H-bonds connect the molecules forming chains along the c-axis. Moreover, there are short intermolecular contacts between the O atoms (3.97)Å and between the O and C atoms (3.98)Å of the nitrophonyl moiety within the chain. Anti-bacterial study reveals that complexes are better anti-microbial agents than three Schiff base due to bacterial cell penetration by chelation.

Keywords : imidazole, pyridine, Crystal Structure, hydrogen bonding.

K. Sakthi Murugesan et al / International Journal of ChemTech Research, 2019,12(4): 277-282.

DOI= http://dx.doi.org/10.20902/IJCTR.2019.120433
