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Studies on complex  $\pi$ - $\pi$  and T-stacking features of imidazole and phenyl/p-halophenyl units in series

of 5-amino-1-(phenyl/p-halophenyl)imidazole-4-carboxamides and their carbonitrile derivatives:

# role of halogens in tuning of conformation

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# **Graphical Abstract**



Studies on complex  $\pi$ - $\pi$  and T-stacking features of imidazole and phenyl/*p*-halophenyl units in series of 5-amino-1-(phenyl/*p*-halophenyl)imidazole-4-carboxamides and their carbonitrile derivatives:

# role of halogens in tuning of conformation

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#### Abstract

5-amino-1-(phenyl/*p*-halophenyl)imidazole-4-carboxamides (N-phenyl AICA) (**2a-e**) and 5-amino-1-(phenyl/*p*-halophenyl)imidazole-4-carbonitriles (N-phenyl AICN) (**3a-e**) had been synthesized. X-ray crystallographic studies of **2a-e** and **3a-e** had been performed to identify any distinct change in stacking patterns in their crystal lattice. Single crystal X-ray diffraction studies of **2a-e** revealed  $\pi$ - $\pi$  stack formations with both imidazole and phenyl/*p*-halophenyl units in *anti* and *syn* parallel-displaced (PD)-type dispositions. No  $\pi$ - $\pi$  stacking of imidazole occurred when the halogen substituent is bromo or iodo;  $\pi$ - $\pi$ stacking in these cases occurred involving phenyl rings only. The presence of an additional T-stacking had been observed in crystal lattices of **3a-e**. Vertical  $\pi$ - $\pi$  stacking distances in *anti*-parallel PD-type arrangements as well as T-stacking distances had shown stacking distances short enough to impart stabilization whereas *syn*-parallel stacking arrangements had got much larger  $\pi$ - $\pi$  stacking distances to belie any *syn*-parallel stacking stabilization. DFT studies had been pursued for quantifying the  $\pi$ - $\pi$ stacking and T-stacking stabilization. The plotted curves for *anti*-parallel and T-stacked moieties had similarities to the 'Morse potential energy curve for diatomic molecule'. The minima of the curves corresponded to the most stable stacking distances and related energy values indicated stacking stabilization. Similar DFT studies on *syn*-parallel systems of **2b** corresponded to no  $\pi$ - $\pi$  stacking stabilization at all. Halogen-halogen interactions had also been observed to stabilize the compounds **2d**, **2e** and **3d**. Nano-structural behaviour of the series of compounds **2a-e** and **3a-e** were thoroughly investigated.

## Keywords

N-phenyl AICA; N-phenyl AICN; X-ray crystallography; C-H<sup> $\dots$ </sup> $\pi$ -stacking; DFT studies, self-aggregation.

# 1. Introduction

Intermolecular interactions are fundamentally important for predicting a gamut of phenomena ranging from simple acid-base behavior to functional nano-material formation [1-6]. In addition to hydrogen bonding interactions,  $\pi$ -  $\pi$  interactions [7], non-classical hydrogen bonds [8-10], halogen-halogen soft interactions [11-16] and non-covalent halogen bonds (X...N/O, X= halogens) [17-18] have been found to be instrumental, in various cases, in determining the physical and chemical properties of compounds with aromatic and heterocyclic rings and halogen substituents. Benzene-dimer serves as the prototypical model of  $\pi$ -  $\pi$  stacking and two most important structural minima for benzene dimers are the parallel-displaced (PD)  $\pi$ -stacked and C-H<sup>...</sup> $\pi$  hydrogen bonded T-stacked structures [19-23]; parameterization of  $\pi$ -  $\pi$ stacking of benzene dimer have been the subject of many theoretical studies involving various functionals in the density functional theory (DFT). In case of parallel-stacked substituted benzene dimers, the aspects of the binding being stronger than the unsubstituted benzene, irrespective of the nature of substituents, have been reported; important contributions of dispersion, induction and exchange-repulsion items to the binding energy have also been considered [21,23,24]. Studies on  $\pi$ -  $\pi$  stacking of aromatically substituted N-heterocyclic molecules have gained momentum [25,26]. Imidazole is an acclaimed heterocyclic nucleus involved in various biochemical processes and pharmaceutics. AICA riboside (1) (Fig. 1) is the de-nevo purine precursor in the bio-synthetic pathway of purine nucleotides [27].



Fig. 1. Structure of AICAR (1)

In continuation of earlier work on AICA [28], recently I have reported [29] a systematic study on the  $\pi$ - $\pi$ stacking features of a series 5-amino-1-alkylimidazole-4-carboxamides. This reports the fact that the imidazole units in these 1-alkylimidazole compounds show, in their X-ray crystallographic patterns, the very important *anti*-parallel and *syn*-parallel  $\pi$ - $\pi$  stackings which were further supported by their DFT analysis. Since gas-phase benzene-dimer has reportedly got a binding energy of 2-3 Kcal mol<sup>-1</sup>[30] I surmised to take up studies on similar AICA compounds where the imidazole unit holds a phenyl/phalophenyl at N-1 instead of alkyl. The aspect of inclusion of an aryl substituent induced possible varieties of  $\pi$ - $\pi$  stackings of homo-dimers (imidazole-imidazole and phenyl-phenyl) as well as hetero-dimer (imidazole-phenyl) types, along with the possible additional intricacies of the presence of anti-parallel and conformations. syn-parallel The conversion from 5-amino-1-(phenyl/p-halophenyl)imidazole-4carboxamides (2a-e) to 5-amino-1-(phenyl/p-halophenyl)imidazole-4-carbonitriles (3a-e) was carried out to study the change in stack formation in the crystal lattice with the change of the functional group. DFT studies on these compounds had been pursued for the purpose of quantifying the stabilization in cases of  $\pi$ - $\pi$  stacking of imidazole dimers and phenyl/p-halophenyl dimers as well as T-stacking interactions. In each case, DFT calculated results were in good agreements with the X-ray crystallographic

results. Recently there has been a surge of studies on  $\pi,\pi$  stacking-based organic materials to unravel unique optical and optoelectronic properties [31] being superior to the corresponding bulk counterparts. 1-D organic nanostructures have been successfully constructed by way of  $\pi$ - $\pi$  stacking, hydrogen bonding [31,32] and electrostatic interactions, among others. Compounds **2a-c**, **2e** and **3a-e** had shown to have generated 1-D nanostructures as well as **2d** had shown nano-vesicle morphology.

# 2. Experimental

#### 2.1. Materials and methods

Melting points were recorded on an electrically heated Köfler Block apparatus and are uncorrected. The solid-state FT-IR measurements were done with a Perkin-Elmer-782 model spectrometer in the form of KBr discs of the compounds. <sup>1</sup>H NMR and <sup>13</sup>C NMR spectra were run with DMSO-d<sub>6</sub> solutions at ambient temperature on a Bruker Avance 300 spectrometer at 300 MHz for <sup>1</sup>H and at 75 MHz for <sup>13</sup>C. Chemical shifts are reported in  $\delta$  and coupling constants in Hz. The multiplicity of the carbon atoms was determined by the DEPT 135° experiment. The DFT calculations had been performed by using the basis set mpwb95/6-31++g for **2a-d** and **3a-d** as well as mpwb95/lanl2dz for **2e** and **3e** in **Gaussian 09W** [33] suite of programs. These basis sets included correlation consistent methods which consist of diffusion correction being advantageous for our targets of  $\pi$ - $\pi$  stacking calculation considered upto long range.

2.2. General procedure for the preparation of 5-amino-1-(phenyl/p-halophenyl) imidazole-4-

# carboxamides (2a-e)

Syntheses of compounds **2a** and **2c** were reported [34]. The same protocol was utilized to synthesize three new compounds **2b**, **2d** and **2e** of the series of **2a-e**.



Scheme 1. Preparation of 5-amino-1-(phenyl/p-halophenyl)imidazole-4-carboxamide

#### compounds (2a-e)

To freshly prepared 2-amino-2-cyanoacetamide (0.02 mol) in dry and freshly distilled acetonitrile (15 ml) triethyl orthoformate (0.02 mol) was added and the mixture was heated under reflux for 40 minutes. To the reaction mixture at room temperature aniline/*p*-haloaniline (0.02 moles) was then added. The reaction mixture was refluxed for 30 min. when the title compound began to precipitate. The reaction mixture was allowed to cool. The ice-cooled mixture was filtered and washed with a little of acetonitrile. The crude product was crystallized from aqueous ethanol.

2.2.1. 5-Amino-1-phenylimidazole-4-carboxamide (2a)

Colourless crystal ( from aq. EtOH), yield 59.4%, m.p. 202 °C: <sup>1</sup>H NMR (300 Mhz, DMSO-d<sub>6</sub>):  $\delta_{\rm H}$  7.37 ( 1H, s, 2-*H*), 5.76 ( 2H, s, O=*C*-N*H*<sub>2</sub>), 6.81 & 6.96 ( 2H, pair of br. s, -N*H*<sub>2</sub>), 7.52 (5H, m, *o*, *m* and *p*-protons); <sup>13</sup>C NMR ( 75MHz, DMSO-d<sub>6</sub>):  $\delta_{\rm C}$  128.57 (2-*C*), 113.48 (4- *C*), 143.09 (5-*C*) , 167.21 (O=*C*-NH<sub>2</sub>), 135.09 (1'-*C*), 125.00 (2'-*C* & 6'-*C*), 130.28 (3'-*C* & 5'-*C*), 130.14 (4'-*C*) ; IR  $\nu_{\rm max}$ (KBr), cm<sup>-1</sup>: 3750 (amide N-H), 3396 (amino N-H), 3189 (aromatic C-H), 1667 (amide –C=O), 1548 (aromatic C=C).

#### 2.2.2. 5-Amino-1-(4-fluorophenyl)imidazole-4-carboxamide (2b)

Colourless crystal ( from aq. EtOH), yield 63.63%, m.p. 282-284°C:  $\delta_{\rm H}$  7.30 (1H, s, 2-*H*), 5.71 ( 2H, s O=C-N*H*<sub>2</sub>), 6.75 & 6.90 ( 2H, pair of br. s, -N*H*<sub>2</sub>), 7.36 (2H, t, J 8.7 Hz, 2'-*H* & 6'-*H*), 7.50 ( 2H, m, 3'-*H* & 5'-*H*), <sup>13</sup>C NMR ( 75MHz, DMSO-d<sub>6</sub>):  $\delta_{\rm C}$  129. 86 (2-*C*), 112.96 (4-*C*), 142.88 (5-*C*), 166.80 (O=*C*-NH<sub>2</sub>), 131.02 (1'-*C*), 127.28 (dd, <sup>3</sup>J<sub>C-F</sub> 9.0 Hz , 2 '-*C* & 6 '-*C*), 116.68 (dd, <sup>2</sup>J<sub>C-F</sub> 23.25 Hz , 3 '-*C* & 5 '-*C*), 161.59 (d, <sup>1</sup>J<sub>C-F</sub> 244.5 Hz ,4'-*C*); IR  $\nu_{max}$ (KBr), cm<sup>-1</sup>: 3770 (amide N-H), 3331.5 (amino N-H), 3110 (aromatic C-H), 1666 (amide –C=O), 1556 (aromatic C=C).

2.2.3. 5-Amino-1-(4-chlorophenyl)imidazole-4-carboxamide (2c)

Colourless crystal ( from aq. EtOH), yield 67.68%, m.p. 258-260°C: <sup>1</sup>H NMR ( 300 Mhz, DMSO-d<sub>6</sub>):  $\delta_{\rm H}$ 7.38( 1H, s, 2-*H*), 5.81 ( 2H, br.*s*, O=C-N*H*<sub>2</sub>), 6.82 and 6.97 ( 2H, pair of br. s., -N*H*<sub>2</sub>), 7.54 (2H, d, J 8.7 Hz, 2'-*H* & 6'-*H*), 7.63 (2H,d, J 8.7 Hz, 3'-*H* & 5'-*H*), <sup>13</sup>C NMR ( 75MHz, DMSO-d<sub>6</sub>):  $\delta_{\rm C}$  130.04 (2-*C*), 113.49 (4-*C*), 143.08 (5-*C*), 167.13 (O=*C*-NH<sub>2</sub>), 133.96 (1'-*C*), 126.95 (2'-*C* & 6'-*C*), 130.18 (3'-*C*)

& 5'-*C*), 133.08 (4'-*C*) ; IR υ<sub>max</sub>(KBr), cm<sup>-1</sup>: 3755 (amide N-H), 3322 (amino N-H), 3103 (aromatic C-H), 1666 (amide –C=O), 1552 (aromatic C=C).

2.2.4. 5-Amino-1-(4-bromophenyl)imidazole-4-carboxamide (2d)

Colourless crystal ( from aq. EtOH), yield 64.05%, m.p. 254-256°C: <sup>1</sup>H NMR ( 300 Mhz, DMSO-d<sub>6</sub>):  $\delta_{\rm H}$ 7.34 (1H, s, 2-*H*), 5.78 ( 2H, br.*s*, O=C-N*H*<sub>2</sub>), 6.78 & 6.94 ( 2H, pair of br. s ,-N*H*<sub>2</sub>), 7.71 (2H, d, J 8.6 Hz, 2'-*H* & 6'-*H*), 7.43 (2H, d, J 8.6 Hz, 3'-*H* & 5'-*H*), <sup>13</sup>C NMR ( 75MHz, DMSO-d<sub>6</sub>):  $\delta_{\rm C}$  129.63 (2-*C*), 113.11 (4- *C*), 142.68 (5-*C*) , 166.76 (O=*C*-NH<sub>2</sub>), 134.00 (1'-*C*), 126.86 (2'-*C* & 6'-*C*), 132.77 (3'-*C* & 5'-*C*), 121.11 (4'-*C*) ; IR  $\nu_{\rm max}$ (KBr), cm<sup>-1</sup>: 3763 (amide N-H), 3332 (amino N-H), 3169 (aromatic C-H), 1655 (amide –C=O), 1540 (aromatic C=C).

2.2.5. 5-Amino-1-(4-iodophenyl)imidazole-4-carboxamide (2e)

Colourless crystal ( from aq. EtOH), yield 61.19%, m.p. 270-272°C: <sup>1</sup>H NMR ( 300 Mhz, DMSO-d<sub>6</sub>):  $\delta_{\rm H}$ 7.32( 1H, s, 2-*H*), 5.76 ( 2H, br.*s*, O=C-N*H*<sub>2</sub>), 6.75 & 6.93 ( 2H, pair of br. s ,-N*H*<sub>2</sub>), 7.27 (2H, d, J 8.4 Hz, 2'-*H* & 6'-*H*), 7.86 (2H, d, J 8.4 Hz, 3'-*H* & 5'-*H*), <sup>13</sup>C NMR ( 75MHz, DMSO-d<sub>6</sub>):  $\delta_{\rm C}$  129.73 (2-*C*), 113.05 (4-*C*), 142.75 (5-*C*) , 166.86 (O=*C*-NH<sub>2</sub>), 134.45 (1'-*C*), 126.95 (2'-*C* & 6'-*C*), 138.75 (3'-*C* & 5'-*C*), 94.10 (4'-*C*) ; IR  $\nu_{\rm max}$ (KBr), cm<sup>-1</sup>: 3763 (amide N-H), 3319 (amino N-H), 3188 (aromatic C-H), 1651 (amide –C=O), 1529 (aromatic C=C) .

2.3. General procedure for the preparation of 5-amino-1-(phenyl/p-halophenyl)imidazole-4-carbonitrils

(**3a-e**)

Synthesis of compound **3c** was reported in literature. Four new compounds **3a-b** and **3d-e** of the series of **3a-e** are reported in this paper.



Scheme 2. Preparation of 5-amino-1-(phenyl/*p*-halophenyl)imidazole-4-carbonitrile compounds (**3a-e**)

5-amino-1-(phenyl/*p*-halophenyl)imidazole-4-carboxamides (2) (4mmol) was added in portions to icecold dimethylformamide-phosphorous oxychloride complex, prepared from DMF (6ml) and POCl<sub>3</sub> (1.2 ml, 12 mmoles), during 15 min with stirring. After the addition was complete, the mixture was kept in the ice bath for 30 min. Then the mixture was allowed to warm up to room temperature and kept at room temperature for a further period of 90 min, stirring being continued all the time. The resulting wine red solution was poured into crushed ice. The solution was made strongly ammoniacal. A brown solid was precipitated. Then it was filtered and the brown mass was warmed on a steam bath with 2% aq. hydrochloric acid (50 ml) for 1 hour. Afterthat, it was allowed to cool in ice-bath and made strongly ammoniacal. The solid was precipitated and ice-cooled mixture was then filtered. The crude product was crystallized from aqueous ethanol.

2.3.1. 5-Amino-1-phenylimidazole-4-carbonitrile (3a)

Colourless crystal ( from aq. EtOH), yield 54.94%, m.p. 182-184 °C: <sup>1</sup>H NMR (300 Mhz, DMSO-d<sub>6</sub>):  $\delta_{\rm H}$ 7.41 (1H, s, 2-*H*), 6.15 (2H, s,-N*H*<sub>2</sub>), 7.45-7.60 (5H, m, *o*, *m* and *p*-protons); <sup>13</sup>C NMR ( 75MHz, DMSO-d<sub>6</sub>):  $\delta_{\rm C}$  133.07 (2-*C*), 91.67 (4-*C*), 147.70 (5-*C*), 117.50 (-*C*N), 134.38 (1'-*C*), 125.77 (2'-*C* & 6'-*C*), 130.34 (3'-*C* & 5'-*C*), 129.17 (4'-*C*); IR(KBr, cm<sup>-1</sup>): 3317 (amino N-H), 3175 (aromatic C-H), 2211(-C=N), 1578 (aromatic C=C).

2.3.2. 5-Amino-1-(4-fluorophenyl)imidazole-4-carbonitrile (3b)

Colourless crystal ( from aq. EtOH), yield 60.10%, m.p. 198-200 °C: <sup>1</sup>H NMR (300 Mhz, DMSO-d<sub>6</sub>):  $\delta_{\rm H}$ 7.45-7.50 (2H, m, 2'-*H* &6'-*H*), 7.33-7.38 (3H, m, 2-*H*, 3'-*H* & 5'-*H*), 6.12 (2H, s, -N*H*<sub>2</sub>); <sup>13</sup>C NMR ( 75MHz, DMSO-d<sub>6</sub>):  $\delta_{\rm C}$  133.09 (2-*C*), 91.39 (4-*C*), 147.96 (5-*C*), 117.47 (-*C*N), 130.64 (1'-*C*, <sup>4</sup>J<sub>C-F</sub> 3 Hz), 128.5 (2'-*C* & 6'-*C*, <sup>3</sup>J<sub>C-F</sub> 9 Hz), 117.11 (3'-*C* & 5'-*C*, <sup>2</sup>J<sub>C-F</sub> 23.25 Hz), 162.31 (4'-*C*, <sup>1</sup>J<sub>C-F</sub> 243.75 Hz).; IR  $\nu_{\rm max}$ (KBr), cm<sup>-1</sup>: 3339.4 (amino N-H), 3191(aromatic C-H), 2368.9, 2217 (-C=N), 1582 (aromatic C=C). 2.3.3. 5-Amino-1-(4-chlorophenyl)imidazole-4-carbonitrile (**3***c*)

Colourless crystal ( from aq. EtOH), yield 55.55%, m.p. 196-198°C: <sup>1</sup>H NMR (300 Mhz, DMSO-d<sub>6</sub>):  $\delta_{\rm H}$ 7.35 (1H, s, 2-*H*), 6.16 (2H, s,-N*H*<sub>2</sub>), 7.44 (2H, d, J 2.1 Hz, J 6.6 Hz, 2'-*H* &6'-*H*), 7.57 (2H, d, J 2.1 Hz, J 6.6 Hz, 3'-*H* & 5'-*H*); <sup>13</sup>C NMR ( 75MHz, DMSO-d<sub>6</sub>):  $\delta_{\rm C}$  132.67 (2-*C*), 91.26 (4-*C*), 147.52 (5-*C*), 117.11 (-*C*N), 133.55 (1'-*C*), 127.55 (2'-*C* & 6'-*C*), 130.00 (3'-*C* & 5'-*C*), 132.88 (4'-*C*); IR  $\upsilon_{\rm max}$ (KBr), cm<sup>-1</sup>: 3324 (amino N-H), 3175 (aromatic C-H), 2213 (-C=N), 1577 (aromatic C=C).

# 2.3.4. 5-Amino-1-(4-bromophenyl)imidazole-4-carbonitrile (3d)

Colourless crystal ( from aq. EtOH), yield 58.82%, m.p. 202-204°C: <sup>1</sup>H NMR (300 Mhz, DMSO-d<sub>6</sub>):  $\delta_{\rm H}$ 7.37 (1H, s, 2-*H*), 6.19 (2H, s,-N*H*<sub>2</sub>), 7.39 (2H, d, J 7.8 Hz, 2'-*H* &6'-*H*), 7.71 (2H, d, J 7.8 Hz, 3'-*H* &5'-*H*); <sup>13</sup>C NMR ( 75MHz, DMSO-d<sub>6</sub>):  $\delta_{\rm C}$  132.47 (2-*C*), 91.19 (4-*C*), 147.36 (5-*C*), 117.03 (-*C*N), 133.26 (1'-*C*), 127.68 (2'-*C* & 6'-*C*), 132.81 (3'-*C* & 5'-*C*), 121.85 (4'-*C*); IR  $\nu_{\rm max}$ (KBr), cm<sup>-1</sup>: 3317 (amino N-H), 3183 (aromatic C-H), 2207 (-C=N), 1568 (aromatic C=C).

2.3.5. 5-Amino-1-(4-iodophenyl)imidazole-4-carbonitrile (3e)

Colourless crystal ( from aq. EtOH), yield 53.19%, m.p. 238-240 °C: <sup>1</sup>H NMR (300 Mhz, DMSO-d<sub>6</sub>):  $\delta_{\rm H}$ 7.40 ( 1H, s, 2-*H*), 6.21 ( 2H, s, -N*H*<sub>2</sub>), 7.28 ( 2H, d, J 8.4 Hz, 2'-*H* & 6'-*H*), 7.91 (2H, d, J 8.4 Hz, 3'-*H* & 5'-*H*); <sup>13</sup>C NMR ( 75MHz, DMSO-d<sub>6</sub>):  $\delta_{\rm C}$  132.83 (2- *C*), 91.64 (4-*C*), 147.67 (5-*C*), 117.42 (-*C*N), 134.11 (1'-*C*), 128.02 (2'-*C* & 6'-*C*), 139.06 (3'-*C* & 5'-*C*), 95.28 (4'-*C*); IR  $\upsilon_{\rm max}$ (KBr), cm<sup>-1</sup>: 3314 (amino N-H), 3184 (aromatic C-H), 2205 (-C=N), 1517.3 (aromatic C=C).

# 2.4. Determination of crystal structure

For the purpose of single crystal X-ray diffraction studies, triple distilled ethanol properly diluted with double distilled water to get aqueous ethanol was used as solvent of crystallization; suitable single crystals of compounds **2a-e** and **3a-e** were obtained by recrystallization of the material from this freshly prepared aqueous ethanol. Diffraction data were collected for **2a-e** and **3a-e** using the freshly crystallized material with MoKα radiation at 296 K using a Bruker APEX-II CCD system. The crystals were positioned 50 mm from the CCD. Frames were measured with a counting time of 10 s. During collection of data none of **2a-e** and **3a-e** showed any evidence of crystal decay. Data analyses were carried out with the Bruker APEX2 and Bruker SAINT programs. The structures were solved using direct methods with the SHELXTL 97 program (Sheldrick, 2008).

The ORTEP diagram of representative compound 5-amino-1-(4-bromophenyl)imidazole-4-carboxamide (2e) of the series 2a-e and 5-amino-1-(4-iodophenyl)imidazole-4-carbonitrile (3d) of the series 3a-e are shown in Fig. 2 and Fig. 3 respectively.





Fig. 2. ORTEP diagram of compound 2e CCDC no. 1050706

Fig. 3. ORTEP diagram of compound 3d CCDC no. 1050710

#### 3. Results and discussion

#### 3.1. Description of crystal structure

The observations that X-ray crystallographic analyses have been successful in unraveling stacked arrangements in case of drug molecules and in the aromatic side-chains of proteins [36, 37] have infused interest in the prediction of the role of stacking interaction on the stabilization of supramolecular architecture. These findings have generated motivation for knowing its role in quantitative structure-activity relationships [38]. The self-assembled supramolecular structure formation is intrigued in many cases by  $\pi$ - $\pi$  stacking as an important non-covalent force. As early as in 1938, Astbury and Bell [39] reported a spacing of 3.3-3.4 Å between flat or almost flat nucleotide units existing perpendicular to the long axis of the molecules. Studies on aromatic stacking interactions using conformationally flexible

molecules have been the focus of a no. of relevant experiments [40]. Although there have been some experimental and DFT studies on the  $\pi$ - $\pi$ -stacking of several systems containing N-heterocyclic units, a recent report [29] of studies on  $\pi$ - $\pi$  stacked imidazole systems have unraveled, for the first time, the role of  $\pi$ - $\pi$  stacking of imidazoles in the *anti*-parallel conformations as the stabilizing factor in the crystal lattice of a series of 5-amino-1-alkylimidazole-4-carboxamides; corresponding syn-parallel conformation failed to tender any appreciable stabilization. I describe here in a systematic way the synthesis of series of compounds 2a-e and 3a-e where the halo moiety embraces all the possible choices out of F, Cl, Br, I. It was expected that the halogens with the widely varying electronegativities and sizes would sometime offer drastically different situations. DFT studies had also been taken up in a cue. Parallel-displaced (PD) [41]  $\pi$ - $\pi$  stack formations had been observed in the X-ray crystallographic studies for **2a-e**, occurring with both imidazole and phenyl/p-halophenyl units in mutually anti-parallel and syn-parallel arrangements. The crystal lattices of **3a-e** revealed the presence of T-stackings along with the PD-stacked *anti-/syn*-parallel  $\pi$ - $\pi$  stackings of both imidazole-imidazole and phenyl-phenyl/p-halophenyl-p-halophenyl rings of two correspondingly adjacent AICA derivatives. The crystal lattices of 2a-e did not show any T-stacking. No hetero-dimer formation was noted in the crystal lattice of 2a-e and 3a-e. In cases of 2d-e and 3a-e some different behaviours were observed.

Table 1 contains the details of crystal data showing the bond lengths and angles of **2a-e** and Table 2 tabulates the similar data of **3a-e**.

Crystal data of	2a-e				
Compound	2a	2b	2c	2d	2e
Formula	C <sub>10</sub> H <sub>10</sub> N <sub>4</sub> O <sub>1</sub>	$C_{10}H_9F_1N_4O_1$	C <sub>10</sub> H <sub>9</sub> Cl <sub>1</sub> N <sub>4</sub> O <sub>1</sub>	C <sub>10</sub> H <sub>9</sub> Br <sub>1</sub> N <sub>4</sub> O <sub>1</sub>	$C_{10}H_9I_1N_4O_1$
M/g mol <sup>-1</sup>	202.22	220.21	236.66	281.12	328.11
Crystal	triclinic	triclinic	triclinic	monoclinic	monoclinic
system					

#### Table 1

Compound	<b>3</b> a	3b	3c	3d 3	le
Table 2     Crystal data of 3	3а-е				
CCDC number	1050702	1050703	1050704	1050705	1050706
Recrys. from	hot aq. ethanol				
wR <sub>2</sub>	0.1707	0.1364	0.1743	0.1190	0.0930
wR <sub>2</sub> R indices (all data) R <sub>1</sub>	0.1561 0.0621	0.1209 0.0470	0.1394 0.0915	0.1060 0.0528	0.0826 0.0522
indices [I>2σ(l)] R <sub>1</sub>					
Final R	0.0533	0.0405	0.0519	0.0380	0.0341
μ / mm-1 Min. T	0.9720	0.9660	0.9016	0.3553	0.4817
Absorption	0.095	0.116	0.353	3.728	2.852
$\rho_{\rm C}$ / g cm <sup>-3</sup>	1.377	1.501	1.524	8 1.701	8 1.952
$V(A^{\circ^3})$	487.83(8)	487.21(14)	515.7(9)	2195.5(8)	2233.5(8)
γ (° )	62.767(4)	63.477(7)	63.289(12)	90.00	90.00
β (° )	87.258(4)	88.210(8)	89.218(13)	134.166(11)	90.462(8)
α (° )	86.642(5)	87.927(8)	85.799(13)	90.00	90.00
c (Å)	9.0756(9)	9.1625(14)	9.366(10)	14.481(3)	15.228(3)
b (Å)	7.9439(7)	7.7666(14)	8.005(8)	10.467(2)	10.732(2)
a (Å)	7.6252(7)	7.6579(12)	7.723(8)	20.193(5)	13.667(3)
Space group	P-1	P-1	P-1	C12/c1	C12/c1

Compound	Ja	50	30	Ju	Se
 Formula	C10H8N4	$C_{10}H_7F_1N_4$	$C_{10}H_7Cl_1N_4$	$C_{10}H_7Br_1N_4$	$\mathrm{C_{10}H_{7}I_{1}N_{4}}$
M/g mol <sup>-1</sup>	184.20	202.20	218.65	263.11	310.10

Crystal system	triclinic	monoclinic	monoclinic	monoclinic	monoclinic
Space group	P-1	C12/c1	P121/c1	P121/c1	P121/c1
a (A°)	6.2970(12)	18.045(5)	18.571(3)	11.1107(7)	10.947(3)
b (A°)	9.248(2)	8.967(2)	18.342(3)	6.4150(4)	6.4268(14)
c (A°)	17.470(3)	12.304(3)	12.0976(17)	15.0571(9)	15.673(4)
α (° )	92.896(10)	90.00	90.00	90.00	90.00
β (° )	91.266(9)	103.385(3)	99.784(2)	105.805(2)	105.432(9)
γ (° )	107.490(10)	90.00	90.00	90.00	90.00
$V(A^{\circ^3})$	968.4(3)	1936.9(9)	4060.9(10)	1032.62(11)	1062.9(4)
Z	4	8	16	4	4
$\rho_C$ / g cm <sup>-3</sup>	1.263	1.387	1.430	1.692	1.938
Absorption $\mu / \text{mm}^{-1}$	0.082	0.103	0.345	3.950	2.983
Correctiontype					
Max. T	0.9919	0.9898	0.9684	0.6486	0.7546
Min. T	0.9758	0.9698	0.9092	0.3646	0.4681
Final Rindices [I>2σ(l)] R <sub>1</sub>	0.0490	0.0471	0.0440	0.0364	0.0276
wR <sub>2</sub> R indices (all data)R <sub>1</sub>	0.1562 0.0671	0.1367 0.0893	0.1328 0.0955	0.0952 0.0508	0.0660 0.0398
wR <sub>2</sub>	0.1843	0.1747	0.1841	0.1055	0.0722
Recrys.from	hot aq. ethanol				
CCDC number	1050707	1050708	1050709	1050710	1050711

The extensive H-bonding patterns of representative compounds 2d of the series 2a-e and 3a of the series 3a-e are shown in Fig. 4. The hydrogen bonding patterns are presented in Supplementary data, Fig. S21

and Fig. S22 as well as the hydrogen bonding parameters for the compounds **2a-e** and **3a-e** are presented in Supplementary data, Table S2 and Table S3.



Fig. 4. a) and b) show hydrogen bonding network of 2d and 3a respectively

The unit cell arrangement of the lattice of representative compound 2c is shown in Fig. 5a. Fig. 5b and Fig. 5c reveal the features of  $\pi$ - $\pi$  stacking of imidazole-imidazole and *p*-chlorophenyl-*p*-chlorophenyl units respectively in PD-type *anti*-parallel arrangements.





**Fig. 5.** (a) unit cell arrangement viewing along b-direction, (b)  $\pi$ - $\pi$  stacking of two *anti*-parallel imidazoles observed in X-ray crystallographic lattice and (c)  $\pi$ - $\pi$  stacking of two *anti*-parallel phenyls observed in X-ray crystallographic lattice of compound **2c**.

Similar *anti*-parallel PD-stacked imidazole-imidazole and phenyl-phenyl  $\pi$ - $\pi$  stacking interactions were observed in the crystal lattices of **2a** and **2b** (shown in Supplementary data, Fig. S54 and Fig. S55 respectively) where the N1 substituents of AICA were phenyl and *p*-fluorophenyl groups respectively. It was further observed from the X-ray studies of **2d** and **2e**, where the halogen substituents were of larger size viz. *p*-bromophenyl- and *p*-iodophenyl- as the N-1 substituents respectively, the closely-associated imidazole rings were not oriented in *anti*-parallel fashion. The closest imidazole rings in the corresponding crystal lattices displayed dihedral angels of 65.49° and 70.41° respectively (shown in Fig. 6a. and Fig. 6b.). These large dihedral angles of closely-associated imidazole rings of **2d** and **2e** exerted difficulty for imidazole-imidazole *anti*-parallel  $\pi$ - $\pi$  stacking.



**Fig. 6.** a) and b) show Dihedral angle between the closest pair of imidazoles in the lattice of **2d** and **2e** respectively

Table 3 presents the *anti*-parallel PD  $\pi$ - $\pi$  stacking distances between imidazole-imidazole of compounds **2a-c** and similar  $\pi$ - $\pi$  stacking distances between phenyl-phenyl moieties of **2a-c** and **2e** with their respective slip angle values. The respective vertical distance values for those compounds are tabulated in Table 3. Another important parameter termed as "vertical distance" [42] is defined as the shortest distance between two stacked molecules and is represented by R<sub>2</sub> (Fig. 7). The value of R<sub>2</sub> could be calculated from the Equation 1 where R<sub>1</sub> is the distances observed between  $\pi$ - $\pi$  stacked parallel-displaced imidazoles and  $\theta$ is designated as slip angle. The slip angle is essentially the angular deviation from the PD-stacked imidazoles and hypothetical sandwich-stacked imidazoles.







Similarly, measurements of slip angle and vertical distances between PD-stacked phenyl/p-halophenyls of compounds 2a-e were calculated (shown in Supplementary data, Fig.S24- Fig.S42). Table 3 shows distances between centroids of two imidazoles and centroids of two phenyls in anti-parallel stacking arrangements were in the range of 4.43-4.53 and 4.01-4.46 Å respectively. These were the experimental anti-parallel PD  $\pi$ - $\pi$  stacking distances. In each case of a PD-stacked dimer, the corresponding vertical distance of *anti*-parallel imidazole dimer as well as that of the phenyl/p-halophenyl dimer are shown in Table 3. The vertical distances in the case of phenyl dimers were in the range of 3.83-4.07 Å and those in the case of imidazole dimers lie in the range of 4.20-4.36 Å (Table 3). Although each of the crystal lattice of 2a, 2b, 2c and 2e corresponded to anti-parallel conformation of the phenyl/p-halophenyl moieties in PD-type arrangements[41], compound 2d had got a sensibly sandwich-type anti-parallel stacking of pbromophenyl moiety. The  $\pi$ - $\pi$  stacking distance can very often be considered as a marker of the strength of stacking interaction. With the results of Ab-initio calculation, Sinnokort et.al. [30] have ascertained a value of 3.4 Å for vertical separation between the planes of PD  $\pi$ - $\pi$  stacked benzenes. The earlier report [29] of the vertical distances of anti-parallel imidazole dimers of 5-amino-1-alkylimidazole-4carboxamides, lying in the range 3.70-3.89 Å, can be compared with  $\pi$ - $\pi$  anti-parallel stacking stabilization.

# Table 3

Compound	imidazole-imid	dazole		Phenyl/p-halop	Phenyl/p-halophenyl-		
	anti-parallel $\pi$ - $\pi$ stacking			phenyl/p-halophenyl			
-		_		anti-parallel $\pi$ - $\pi$	t stacking		
	$\pi$ - $\pi$ stacking	slip angle	vertical	$\pi$ - $\pi$ stacking	slip angle	vertical	
	distance	from	π-π	distance from	from	π-π	
	from crystal	crystal	distance	crystal	crystal	distance	
	lattice (Å)	lattice (°)	(Å)	lattice(Å)	lattice(°)	(Å)	

Anti-parallel PD  $\pi$ - $\pi$  stacking distances from X-ray crystallographic analysis of **2a-e** and the corresponding vertical  $\pi$ - $\pi$  stacking distances

2a	4.53	16.08	4.35	4.14	15.19	3.99
2b	4.43	18.30	4.20	4.46	30.88	3.83
2c	4.53	15.63	4.36	4.01	6.49	3.98
2d	-	-	-	<b>4.04</b> <sup>#</sup>	- 6	-
2e	-	-	-	4.20	14.15	4.07

# PD  $\pi$ - $\pi$  stacking distance between *p*-bromophenyls for **1d** is essentially "Sandwich" stacking A comparison of vertical anti-parallel phenyl-phenyl stacking distances (Table 3) also revealed that vertical distance for fluoro compound (2b) is the least (3.83 Å) and this value increased steadily in going from fluoro to iodo in the series of 2b-2e. Among all the halogens, fluorine with the highest electronegativity value of 4.0, a negative inductive effect value ( $\sigma_I$ ) of 0.51 and a positive mesomeric effect ( $\sigma_R$ ) of -0.34 can show strong intramolecular interactions in case of organic fluoro compounds [43]. In 2000, Prasanna and Guru Row [44] have published a data base study on the influence of C-F... $\pi$ interactions on crystal packing and conformations. Recently S.Terada et.al. [45] have published an account of polymorphism of different fluoro-substituted aromatic sulfonamides showing dependence on intermolecular  $\pi$ - $\pi$  interactions. Sinnokrot *et. al.* [24] have earlier communicated DFT studies on the effect of substituent of a substituted-benzene unit on the binding with an unsubstituted benzene where -F and -CN substituents correspond to much higher binding energies and lower distances between the centroids of the rings. It had been discussed earlier that the anti-parallel PD  $\pi$ - $\pi$  stackings between imidazoleimidazole pairs were not feasible in cases of 2d-e. The phenyl-phenyl stacking for the compound 2d was actually "Sandwich" (S) in nature which was evidenced from Fig. 8b. Anti-parallel PD-stacked piodophenyl-p-iodophenyl  $\pi$ - $\pi$  stacking interaction of compound 2e is shown in Supplementary data, Fig.S56.



**Fig. 8.** (a) unit cell arrangement viewing along b-direction of **2d**, (b) sandwich type  $\pi$ - $\pi$  stacking of two *anti*-parallel *p*-bromophenyls

Literature survey revealed that PD-type  $\pi$ - $\pi$  stacking was generally more stabilizing in nature than the Stype [30]. Overall stabilization of the lattices of **2d** and **2e** must have come from a balance of other stabilizing features e.g. halogen-halogen soft interactions etc. Simultaneously, we could take out the closest *syn*-parallel dimer in the same fashion to get the  $\pi$ - $\pi$  stacking distances between the centroids of the two imidazole units and the centroids of the two phenyl units to have the identical values, irrespective of slightly different slip angle [42]. They lie in the range of 7.66-11.63 Å. Table 4 shows the *syn*-parallel  $\pi$ - $\pi$ stacking distances between imidazole-imidazole and phenyl-phenyl stacked moieties obtained from X-ray crystallographic analysis for the compounds **2a-e**. Due to variation in the slip angles between the  $\pi$ - $\pi$ stacked imidazole-imidazole and phenyl-phenyl moieties, differences in vertical distances for those pairs were observed.

#### Table 4

Syn-parallel  $\pi$ - $\pi$  stacking distances from X-ray crystallographic analysis of **2a-e** and the corresponding vertical  $\pi$ - $\pi$  stacking distances

Compound	imidazole-imi	imidazole-imidazole			Phenyl/p-halophenyl-		
	anti-parallel $\pi$	-π stacking		phenyl/p-halop	ohenyl		
				anti-parallel π-	$\pi$ stacking		
	$\pi$ - $\pi$ stacking	slip angle	vertical	$\pi$ - $\pi$ stacking	slip angle	vertical	
	distance	from	π-π	distance	from crystal	π-π	
	from crystal	crystal	distance	from crystal	lattice(°)	distance	
	lattice	lattice (°)	(Å)	lattice(Å)		(Å)	
	(Å)						

2a	7.94	30.91	6.81	7.94	25.45	7.17
2b	7.66	21.07	7.14	7.66	26.20	6.87
2c	8.25	55.24	4.70	8.25	54.69	4.77
2d	11.63	17.62	11.08	11.63	19.31	10.97
2e	8.69	12.79	8.47	8.69	13.63	8.44

It was interesting to note that, for compounds **2a-e**, the phenyl/*p*-halophenyl units are attached to respective N-1 of imidazole units with a free-rotable single bond. The lattices showed that the values of torsion angles between the imidazole and the respective N1 phenyl/*p*-halophenyl ring for those AICA derivatives were in the range of  $|43.75^{\circ}|$  to  $|52.16^{\circ}|$  (please see Supplementary data, Table S4). As a consequence, mixed imidazole-phenyl stacking was not possible in crystal lattices of **2a-e**. Occurrences of preferential intramolecular  $\pi$ - $\pi$  stabilizing interactions between two aromatic components of a sufficiently flexible molecule is documented [46-50]. These molecules are known as foldamers [51]. This phenomenon can be related to the optimized conformation of a molecule in its crystalline lattices maintaining a balance of size, electronegativity and electronaffinity of the halogens/substituents (please see Supplementary data, Table S1). None of the molecules **2a-e** exhibited any folding characteristics.

Then the attention was drawn towards the possible mode of  $\pi$ -  $\pi$  stackings in the compounds **3a-e** where the amide groups at C-4 position of imidazole rings of each of the AICA derivatives **2a-e** was converted to cyano groups keeping phenyl/*p*-halophenyl substituents as N1 of imidazole moiety intact (Scheme 2). The crystal lattices of **3a-e** revealed the presence of T-stackings along with the *anti* /*syn*  $\pi$ - $\pi$  stackings of both imidazoles and phenyl/*p*-halophenyl moiety. The unit cells of the lattices of two representative compounds **3a** and **3b**, their T-stacking patterns and *anti*-parallel  $\pi$ - $\pi$  stackings of imidazole-imidazole and phenyl-phenyl moieties are shown in Fig. 9 and Fig. 10 respectively.



**Fig. 9.** (a) unit cell arrangement viewing along b-direction, (b) T-stacking between the hydrogen attached with the C-2 position of imidazole and the closest phenyl moiety, (c) T-stacking between the hydrogen attached with the C-4' position of phenyl and the closest phenyl moiety and (d)  $\pi$ - $\pi$  stacking distance of two *anti*-parallel imidazole-imidazole of compound **3a** 



**Fig. 10.** (a) unit cell arrangement viewing along b-direction, (b) T-stacking distance between the hydrogen attached with the C-2 position of imidazole and the closest *p*-fluorophenyl moiety, (c)  $\pi$ - $\pi$  stacking distance of two *anti*-parallel imidazoles and (d)  $\pi$ - $\pi$  stacking distance of two *anti*-parallel *p*-fluorophenyls of compound **3b**.

X-ray crystallographic studies of **3a-e** indicated the presence of imidazole-imidazole PD  $\pi$ - $\pi$  stacking in *anti*-parallel fashion (Fig. 9d for compound **3a** and Fig. 10c for compound **3b**) in their crystal lattices. *Anti*-parallel phenyl-phenyl PD  $\pi$ - $\pi$  stackings were observed in the crystal lattices of **3b-e** (Fig. 10d for compound **3b**) whereas similar stacking pattern was not detected in the crystal lattice of **3a**. It was fascinating to note that **3a** showed its differences with **3b-e** in their T-stacking pattern. The crystal lattice of **3a** had shown dual T-stacking interactions. Each of the molecules belonging to the series **3a-e** had specific T-stacking between C2-H of imidazole ring with the closest  $\pi$ -cloud of phenyl moiety (Fig. 9a and Fig. 10a). The crystal lattice of **3a** disclosed additional T-stacking involving C4'-H of N1-phenyl substituent with the closest  $\pi$ -cloud of phenyl moiety (Fig. 9b). This type of extra T-stackings were not identified in the crystal lattices of **3b-e**, since in those molecules the C-4' positions of phenyl substituents at N1 of their imidazole rings were occupied by -F, -Cl, -Br and -I respectively. T-stacking between C2-H of imidazole ring caused by the electron-withdrawing effect of –CN group at C4 position. T-stacking, *anti*-parallel PD-stacked imidazole-imidazole and phenyl-phenyl  $\pi$ - $\pi$  stacking interactions of compounds **3c-e** are presented in Supplementary data, Fig.S57-Fig.S59.

# Table 5

Compound	imidazole-imidazole anti-parallel $\pi$ - $\pi$ stacking			phenyl-phenyl $anti-parallel \pi-\pi$ stacking			
	$\pi$ - $\pi$ stacking distance from X-ray crystallogra- phic lattice(Å)	slip angle from X-ray crystallog- raphic lattice(°)	vertical π-π distance (Å)	$\pi$ - $\pi$ stacking distance from X-ray crystallog- raphic lattice(Å)	slip angle from X-ray crystallogr ap-hic lattice(°)	vertical π-π distance (Å)	
3a 3b	4.61 4.09	4.58 5.60	4.59 4.07	4.19	35.18	- 3.42	

Anti-parallel PD $\pi$ - $\pi$ stacking distances from 2	X-ray crystallographic analysis of <b>3a-e</b> and the
corresponding vertical $\pi$ - $\pi$ stacking distances	

3c	4.92	7.45	4.88	4.88	43.24	3.55
3d	4.18	36.64	3.35	4.05	7.18	4.12
<b>3e</b>	4.03	32.07	3.41	4.16	7.71	4.12

Table 5 tabulates the distances between the centroids of the two closest imidazoles in *anti*-parallel stacking arrangements of **3a-e** and also the distances between PD-stacked *p*-halophenyl–*p*-halophenyl moieties for **3b-e**. Corresponding vertical distances values for each of the PD-stacked imidazole pairs and *p*-halophenyl pairs are reported in Table 5 along with the characteristic slip-angles (shown in Supplementary data, Fig.S43-Fig.S53). The vertical distances which are important for the measurements of the closeness of two PD-stacked moieties are not applicable in T-stacking interactions which involve C-H<sup>---</sup> $\pi$  type interactions instead of PD-stacked  $\pi$ - $\pi$  type. Simultaneously, the closest *syn*-parallel dimers from the crystal lattices of **3a-e** were taken out to obtain the  $\pi$ - $\pi$  stacking distances between the centroids of the two PD-stacked imidazole units and between the centroids of the two phenyl units. As observed in cases of **2a-e**, both types of PD-stacked arrangements for **3a-e** showed identical distance values in respective cases with explicit differences in slip angles (shown in Table 6). Since the phenyl/*p*-halophenyl unit is attached to N-1 of imidazole with a free-rotable single bond, the lattices showed preference for the torsion angles between imidazole and phenyls to lie in the range of  $|50.66^{\circ}|$  to  $|78.36^{\circ}|$  (please see Supplementary data, Table S5) in cases of **3a-e**. Hetero imidazole-phenyl  $\pi$ - $\pi$  stacking was not possible also in the molecules **3a-e**.

#### Table 6

	0						
Compound	imidazole-imi	imidazole-imidazole			phenyl-phenyl		
	syn-parallel π-	-π stacking		syn-parallel $\pi$ -	$\pi$ stacking		
	$\pi$ - $\pi$ stacking	slip angle	Vertical	$\pi$ - $\pi$ stacking	slip angle	Vertical	
	distance	from X-	π-π	distance	from X-ray	π-π	
	from X-ray	ray	distance	from X-ray	crystallogr-	dista-	
	crystallogr- aphic	graphic lattice(°)	(Å)	crystallog- raphic	lattice(°)	lice (A)	
	lattice(Å)	()		lattice(Å)			

Syn-parallel $\pi$ - $\pi$ stacking distances from X-ra	y crystallographic analysis	of 3a-e and the	corresponding
vertical $\pi$ - $\pi$ stacking distances			

3a	6.23	46.50	4.29	6.23	40.93	4.71	
3b	6.18	38.22	4.85	6.29	33.97	5.22	
3c	6.08	38.88	4.73	6.29	30.42	5.42	
3d	6.41	44.03	4.61	6.41	36.33	5.16	
<b>3e</b>	6.43	44.16	4.61	6.43	36.92	5.14	

Table 7 presents T-stacking distances involving C2-H of imidazaole and the closest  $\pi$ -cloud of phenyl/*p*-halophenyl moieties in the crysyal lattices of **3a-e**. The values were in the range of 2.92-3.16Å and might afford stability to these systems. Another T-stacking was recognized in the crystal lattice of **3a** involving C4'-H of phenyl substituent and the closest  $\pi$ -cloud of phenyl moiety (Fig. 9c) and presented in Table 7.

#### Table 7

T- stacking distances from X-ray crystallographic analysis of 3a-e

Compound	T-stacking distance	T-stacking stacking
	between C-2 hydrogen	between C-4'hydrogen
	of imidazole and the closest	of phenyl and the closest
	phenyl/p-halophenyl moiety (Å)	phenyl ring (Å)
3a	2.96	3.00, 3.01
3b	2.98	-
3c	2.92	-
3d	3.16	-
3e	3.15	-

T-stacking distances of **3a-e** were the shortest stacking distances than the distances corresponding to *anti*parallel imidazole-imidazole and phenyl-phenyl stacked moities whereas the *syn* parallel PD-stackings for those molecules recorded the highest values.

# 3.2. Computational studies on stacking interactions

At an early date, rotational contour analysis of the electronic spectra of *sym*-tetrazine has revealed the presence of two isomers: a planar C-H<sup>...</sup>N hydrogen bonded dimer and a T-shaped dimer with C-H<sup>...</sup> $\pi$  interactions [52]. Using a combination of two-colour resonant two-photon ionization technique and

Leonard-Jones potential calculation [53], both syn-parallel and anti-parallel  $\pi$ -stacked and T-stacked varieties of pyrazine-dimers have been reported [52]. As evident from the order of stability of dimers as 1,3,5-triazine> pyrazine> pyridine, reported by Sathamurthy et. al. [54], an increase of the number of nitrogen atoms on the heterocyclic ring leads to an increased  $\pi$ -stacking strength. A comparison of energetics of benzene-dimer, benzene-pyridine hetero-dimer and pyridine-dimer with the help of CCSD(T) level of calculation indicates that the introduction of nitrogen to replace the carbon atom of benzene creates a dipole along with reduction of spatial extent of  $\pi$  cloud and polariziability of heteroaromatic moiety comparative to benzene [52,54,26]; the stabilization in the case of pyridine dimer has shown to be of the order of 1.43 Kcal mol<sup>-1</sup> higher than the benzene dimer [26,54]. It is also to note that the results of *ab-initio* calculations on the benzene-dimers depend on the selection of basis set to some extent [42,55]. A detailed understanding of the extent of stabilization offered by different  $\pi$ - $\pi$  stacking and T-stacking interactions of compounds 2a-e and 3a-e was correlated with the results of DFT calculations. For compounds 2a-e, DFT studies were focused on PD-stacked (a) anti/syn-parallel imidazole pairs and (b) anti/syn-parallel phenyl pairs occurring simultaneously in their crystal lattices. For the consideration of anti-parallel stackings the studies had focused on the closest pairs of such anti-parallel imidazoles in void and also such closest pair of anti-parallel phenyls/p-halophenyl in void, in the respective cases, by way of proper selection out of X-ray crystallographic packing keeping the mode of  $\pi$ - $\pi$  stacking as well as the slip angle intact (Table 2 and Table 3). This would afford the optimized form of the respective dimer at once. In a cue, detailed DFT calculations of the optimized energies of dimers using the basis set mpwb95/6-31++g for 2a-d and mpwb95/lanl2dz [56] for 2e had been pursued by keeping the slip-angle of  $\pi$ - $\pi$ stacked dimers unaltered while adopting gradual change of  $\pi$ - $\pi$  stacking distance parameter. Subsequently, the  $\pi$ - $\pi$ -stacking energy corresponding to each set of calculations for a given  $\pi$ - $\pi$  distance is computed from the Equation 2.

 $\pi$ - $\pi$  stacking energy= (optimized energy of the dimer) - 2x (optimized energy of the

# monomer).....Equation 2

The optimized energy of the monomer, remaining the same for the anti-parallel conformations of imidazoles as well as phenyl/p-halophenyl, had been separately calculated by using the same basis sets mpwb95/6-31++g for 2a-d and mpwb95/lanl2dz for 2e. The crystallographic CIF was used to generate the initial geometery of the stacked dimer and monomer. Afterthat, single point energy had been calculated using DFT method. Here 'optimized geometry' indicates actually the geometry which had already been optimized by X-ray crystallographic analysis in the solid state. This calculated  $\pi$ - $\pi$  stacking energy corresponded to simple  $\pi$ - $\pi$  stacking situation only, infact unmixed with other possible interactions like hydrogen bonding. Here mpwb95/6-31++g and mpwb95/lanl2dz basis sets were used since these correspond to correlation consistent methods [56,57] and included the features of diffusion correction as was appropriate for these types of  $\pi$ - $\pi$  stacking calculations considered up to a long stacking distance (total data of the calculations have been tabulated in Supplementary data, Table S6-S13). A graph was drawn by plotting the  $\pi$ - $\pi$ -stabilization energy data against variable  $\pi$ - $\pi$ -stacking distance. DFT calculations on  $\pi$ - $\pi$  stacking of representative compounds 2c and 2e are shown in Fig. 11 and Fig. 12 (DFT calculations on  $\pi$ - $\pi$  stacking of compounds 2a, 2b and 2d are shown in Supplementary data, Fig. S60-Fig.S62). Fig. 11b represents  $\pi$ - $\pi$ -stacking distance of two imidazoles in *anti*-parallel orientation of representative compound 2c as appeared in its crystal lattice. Fig. 11c shows DFT-generated curve of  $\pi$ - $\pi$ stacking energies vs. variable  $\pi$ - $\pi$  stacking distances as computed for the compound 2c. Similarly for corresponding *anti*-parallel phenyl/p-halophenyl stacking, Fig. 11d and Fig. 12b exhibit  $\pi$ - $\pi$  -stacking distances obtained from the crystal lattices of 2c and 2e respectively and Fig. 11e and Fig. 12c indicate DFT-generated curve of compound 2c and 2e respectively.



**Fig. 11.** (a) unit cell arrangement viewing along b-direction, (b)  $\pi$ - $\pi$  stacking distance of two *anti*-parallel imidazoles, (c) the plot of  $\pi$ - $\pi$  stacking distance of *anti*-parallel imidazoles, (d)  $\pi$ - $\pi$  stacking distance of two *anti*-parallel *p*-chlorophenyls and (e) the plot of  $\pi$ - $\pi$  stacking distance of two *anti*-parallel *p*-chlorophenyls and (e) the plot of  $\pi$ - $\pi$  stacking distance of two *anti*-parallel *p*-chlorophenyls and (e) the plot of  $\pi$ - $\pi$  stacking distance of two *anti*-parallel *p*-chlorophenyls and (e) the plot of  $\pi$ - $\pi$  stacking distance of two *anti*-parallel *p*-chlorophenyls and (e) the plot of  $\pi$ - $\pi$  stacking distance of two *anti*-parallel *p*-chlorophenyls, corresponding to compound **2c** 



**Fig. 12.** (a) unit cell arrangement viewing along b-direction, (b)  $\pi$ - $\pi$  stacking distance of two *anti*-parallel *p*-iodophenyls and (c) the plot of  $\pi$ - $\pi$  stacking distance of two *anti*-parallel *p*-iodophenyls, corresponding to compound **2e** 

Thus maximum stacking stabilization energies corresponding to the minimum  $\pi$ - $\pi$ -stacking distances are known from the curve generated by DFT calculations. Table 8 furnishes an effective comparison of *anti*-parallel  $\pi$ - $\pi$  stacking distances of imidazoles pairs of **2a-c** with those deduced from DFT calculations.

Table 9 tabulates similar comparisons for *anti*-parallel  $\pi$ - $\pi$  stacking distance of phenyls/p-halophenyls of

# 2а-е.

# Table 8

Comparison of PD  $\pi$ - $\pi$  stacking distances for the *anti*-parallel pairs of imidazoles of **2a-c** obtained from X-ray crystallographic and DFT studies

Compound	π-π stacking distance(Å) from X-ray crystallographic analysis	min. π-π stacking distance(Å) obtained from DFT calculations	stacking energy (Kcal-mol <sup>-1</sup> ) from DFT calculations
2a	4.53	4.3	-12.80
2b	4.43	4.2	-11.94
2c	4.53	4.2	-23.80

# Table 9

Comparison of PD  $\pi$ - $\pi$  stacking distances for the *anti*-parallel pairs of phenyls of **2a-e** obtained from X-ray crystallographic and DFT studies

compound	π-π stacking distance(Å) from X-ray crystallographic analysis	min. π-π stacking distance(Å) obtained from DFT calculations	stacking energy (Kcal-mol <sup>-1</sup> ) from DFT calculations
2a	4.14	4.2	-2.90
<b>2b</b>	4.46	4.4	-3.84
2c	4.01	4.1	-1.54
2d	4.04	3.8	-12.81
2e	4.2	4.3	-2.31

Comparisons of the data, presented in Table 8 and Table 9, reveal that stacking energies for *anti*-parallel  $\pi$ - $\pi$ -stackings of imidazole pairs are in the range of 11.94 to 23.80 Kcal mol<sup>-1</sup> whereas the *anti*-parallel  $\pi$ - $\pi$  stacking energies of phenyl/*p*-halophenyl pair are much less viz.1.54-3.84 Kcal mol<sup>-1</sup> with the exception

of *p*-bromophenyl-AICA (2d) having an oddly high value of 12.81 Kcal mol<sup>-1</sup>. Studies on *syn*-parallel stackings of respective imidazole pairs and phenyls/*p*-halophenyl pairs were also undertaken. Compound 2b possessing fluorine at the 4'-position of aromatic ring was chosen. The closest pair of imidazoles and that of *p*-fluorophenyls are taken out from the crystal lattice (shown in Fig. 13a) as the optimized pairs (Fig. 13b). As discussed earlier,  $\pi$ -  $\pi$  stacking distances for both *syn*-parallel imidazole pair and *syn*-parallel *p*-fluorophenyl are the same and the value is 7.66Å. DFT calculations were performed keeping slip-angle description unaltered while adopting gradual change of  $\pi$ -  $\pi$  stacking distance parameter. Afterwards, the apparent  $\pi$ - $\pi$  stacking energies were computed from the Equation 2 and plotted against  $\pi$ - $\pi$  stacking distances to get the curve (Fig. 13c).



**Fig. 13.** (a) unit cell arrangement viewing along a-direction, (b)  $\pi$ - $\pi$  stacking distances of two *syn*-parallel imidazoles and *syn*-parallel *p*-fluorophenyls in the lattice and (c) the plot of  $\pi$ - $\pi$  stacking energies against varying  $\pi$ - $\pi$  stacking distance of *syn*-parallel dimer, corresponding to compound **2b** (slip angle kept intact as observed in X-ray crystallographic lattice).

It should be mentioned here that for *syn*-parallel stackings, not only the respective imidazole pairs and *p*-fluorophenyl pairs reinforce their stacking mutually, hydrogen bonding and other weak forces also contribute to the stability of *syn*-parallel dimer system. The graph would be of an exponential nature and can be attempted for fitting with the use of gnuplot; taking the help of the following three possible functions (errors quoted are '1 $\sigma$ ' errors).

(ii)  $F2(x) = \exp(a x + b)$  with best fit values where a = -3.44684 + -0.1632 (4.736%), b = 25.5005 + -0.9679 (3.796%) and reduced chi-square = 9.42956,

(iii) F3(x) = c + exp (a x + b) with best fit values where a = -3.14733 + -0.07875(2.502%), b = 23.7276 + -0.4639 (1.955%), c = -3.80906 + -0.6201 (16.28%) and reduced chi-square = 1.74827

The asymptotic part of the curve starts from a PD  $\pi$ - $\pi$  distance of 7.6 Å and this appeared to be almost same with the  $\pi$ - $\pi$  stacking distance of 7.66 Å observed in the crystal lattice of **2b**. The graphically calculated  $\pi$ - $\pi$  stacking energy was -2.42Kcal mol<sup>-1</sup> which included all types of non-covalent interactions. To ascertain the similar correlation between X-ray crystallographic data and DFT calculated results in case of 3a-e, the studies were focused on (a) T-stacking between C2-H of imidazole and the closest phenyl/p-halophenyl moiety for **3a-e** with an additional T-stacking between C4'-H of imidazole and the closest phenyl moiety in 3a as well as (b)  $\pi$ - $\pi$  stacking between parallel-displaced *anti* -parallel imidazole and (c) anti -parallel phenyl pairs involving gradual changes of C-H<sup> $\dots$ </sup> $\pi$  and  $\pi$ - $\pi$  stacking distance parameters respectively. The optimized energy of the monomer of 3a-e was calculated by using the basis set mpwb95/6-31++g for **3a-d** and mpwb95/lanl2dz for **3e.** DFT calculations on T-stacking and  $\pi$ - $\pi$  stacking of representative compounds 3a and 3c are shown in Fig. 14 and Fig. 15 (DFT calculations on T-stacking and  $\pi$ - $\pi$  stacking of compounds 3b, 3d and 3e are shown in Supplementary materials, Fig. S63-Fig.S65). Fig. 14b and Fig. 15b show the T-stacking distances between C2-H of imidazole and the closest phenyl/p-halophenyl moiety of representative compounds 3a and 3c respectively while Fig. 14d indicates the T-stacking distance between C4'-H of phenyl and the closest phenyl moiety of **3a** as revealed from their respective crystal lattices and corresponding Fig. 14c, Fig. 14e and Fig. 15c specify the curves of T-stacking energy vs. variable T -stacking distance obtained from the DFT calculation. Fig. 14f and

Fig. 15d show  $\pi$ - $\pi$ -stacking distances of PD-stacked *anti*-parallel imidazole pairs and corresponding Fig. 14g and Fig. 15e indicate the relevant curves for DFT calculated  $\pi$ - $\pi$ -stacking energies *vs*. variable  $\pi$ - $\pi$ -stacking distances for compounds **3a** and **3c** respectively. Fig. 15f gives  $\pi$ - $\pi$ -stacking distance for *anti*-parallel *p*-chlorophenyl pairs in the crystal lattice of **3c** and Fig. 15g shows the respective DFT-generated curve for the variables as stated in all previous cases.



**Fig. 14.** (a) unit cell arrangement viewing along b-direction, (b) T-stacking distance between the hydrogen attached with the C-2 position of imidazole and the closest phenyl moiety, (c) the plot of the T-stacking energies against varying T-stacking distances between the hydrogen attached with the C-2 position of imidazole and the closest phenyl moiety, (d) T-stacking distance between the hydrogen attached with the C-4' position of phenyl and the closest phenyl moiety, (e) the plot of T-stacking energies against varying T-stacking distances between the hydrogen attached with the C-4' position of phenyl and the closest phenyl moiety, (e) the plot of T-stacking energies against varying T-stacking distances between the hydrogen attached with the C-4' position of phenyl and the closest phenyl moiety, (f)  $\pi$ -  $\pi$  stacking distance of two *anti*-parallel imidazoles and (g) the plot of the  $\pi$ -  $\pi$  stacking energies against varying  $\pi$ -  $\pi$  stacking distance of two *anti*-parallel imidazoles, corresponding to compound **3a** 



**Fig. 15.** (a) unit cell arrangement viewing along b-direction, (b) T-stacking distance between the hydrogen attached with the C-2 position of imidazole and the closest *p*-chlorophenyl moiety, (c) the plot of the T-stacking energies against varying T-stacking distances between the hydrogen attached with the C-2 position of imidazole and the closest *p*-chlorophenyl moiety, (d)  $\pi$ - $\pi$  stacking distance of two *anti*-parallel imidazoles, (e) the plot of the  $\pi$ - $\pi$  stacking energies against varying  $\pi$ - $\pi$  stacking distances of two *anti*-parallel imidazoles, (f)  $\pi$ - $\pi$  stacking distance of two *anti*-parallel *p*-chlorophenyls and (g) the plot of the  $\pi$ - $\pi$  stacking energies against varying  $\pi$ - $\pi$  stacking distances of two *anti*-parallel *p*-chlorophenyls, corresponding to compound **3c** 

Total data of the DFT calculations of compounds **3a-e** have been tabulated in Supplementary data, Table S15-S29). Table 10 displays all the data relating to both types of T-stackings observed in the molecules **3a-e** including the specific one manifested in case of **3a** which was absent in **3b-e**.

# Table 10

Comparison of T-stacking distances between C2-H of imidazole and the closest phenyl/*p*-halophenyl moieties obtained from X-ray crystallographic and DFT studies of **2a-e** 

Compound	T-stacking between C2-H of imidazole and the closest phenyl/ <i>p</i> -halophenyl moities		T-stacking between C4'-H of imidazole and the closest phenyl/p- halophenyl moities			
	T-stacking distance(Å) from X-ray crystallograp -hic analysis	T-stacking distance (Å) obtained from DFT calculation	Stacking energy (Kcal- mol <sup>-1</sup> ) from DFT calculation	T- stacking distance (Å) from X-ray crystall- ographic analysis	T-stacking distance (Å) obtained from DFT calculation	stacking energy (Kcal- mol <sup>-1</sup> ) from DFT calculation
2a	2.96	3.0	-40.58	3.00, 3.01	3.2, 3.5	-16.70, -16.71
2b	2.98	2.9	- 7.56	<u>-</u>	-	-
2c	2.92	2.8	-7.64	-	-	-
2d	3.16	3.1	-10.53	-	-	-
2e	3.15	3.1	-8.04	-	-	-

A comparison between the observed  $\pi$ - $\pi$  stacking distances for *anti*-parallel pairs of imidazoles and related DFT computational data for the compounds **3a-e** is presented in Table 11. Table 12 tabulates analogous comparison for *anti*-parallel  $\pi$ - $\pi$  stacking involving phenyls/*p*-halophenyls pairs of compounds **3b-e**.

# Table 11

Comparison of PD  $\pi$ - $\pi$  stacking distances of the *anti*-parallel pairs of imidazoles obtained from X-ray crystallographic and DFT studies of **3a-e** 

Compound	<ul> <li>π-π stacking distance(Å)</li> <li>from X-ray</li> <li>crystallographic analysis</li> </ul>	π-π stacking distance(Å) obtained from DFT calculation	stackingenergy (Kcal-mol <sup>-1</sup> ) from DFT calculation
3a	4.61	4.3	-55.73
<b>3</b> b	4.09	3.75	-11.50
<b>3</b> c	4.92	4.7	-0.92
<b>3d</b>	4.18	4.1	-9.54
<b>3e</b>	4.03	3.9	-9.60

# Table 12

Comparison of PD  $\pi$ - $\pi$  stacking distances for the *anti*-parallel pairs of phenyls/*p*-halophenyls obtained from X-ray crystallographic and DFT studies of **3b-e** 

Compound	<ul> <li>π-π stacking</li> <li>distance(Å)</li> <li>from X-ray</li> <li>crystallographic</li> <li>analysis</li> </ul>	<ul> <li>π-π stacking distance(Å)</li> <li>obtained from</li> <li>DFTcalculation</li> </ul>	stackingenergy (Kcal-mol <sup>-1</sup> ) from DFT calculation
<b>3</b> b	4.19	4.3,4.7	-1.25, -1.26
3c	4.88	5.2	-6.07
3d	4.05	3.8	-9.94
<b>3</b> e	4.16	4.15	-1.62

3.3. Skew-syn and skew-anti arrangements of 2b and 2c vis-à-vis computational study

Compounds **3a**, **3d** and **3e** had shown normal *anti* and *syn* parallel arrangements in their crystal lattices where dihedral angels were 0° and 180° for *syn* and *anti* arrangements (please see Supplementary data, Fig.S66 and Fig.S67). However, the compounds **3b** and **3c** with flourine and chlorine respectively at 4'-

position of the aromatic substituent on N1-imidazole ring had shown different behaviours with respect to other derivatives. In their crystal lattices, the  $\pi$ - $\pi$  stacked phenyl rings did not make 0° and 180° dihedral angles in their *syn*-parallel and anti-parallel arrangements. Hence instead of describing them as *syn* and *anti* nomenclatures, arrangements in crystal lattices of **3b** and **3c** were categorized as skew-*anti* and skew-*syn*. Fig. 16 and Fig. 17 show skew-*syn* and skew-*anti* orientation in the crystal lattices of **3b** and **3c** respectively.



Fig. 16. a) and b) show skew-syn and skew-anti orientation respectively in the crystal lattice of 3b



**Fig. 17.** a) and b) show skew-*syn* and skew-*anti* orientation respectively in the crystal lattice of **3c** X-ray crystallographic studies had shown that the dihedral angels of skew-*syn* and skew-*anti* orientation of **3b** were 29.25° and 149.54° (Fig. 18) respectively and the dihedral angles of skew-*syn* and skew-*anti* orientation of **3c** were 41.27° and 136.35° respectively (Fig.19).



Fig. 18. a) and b) show dihedral angles of skew-syn and skew-anti orientation of 3b



Fig. 19. a) and b) show dihedral angles of skew-syn and skew-anti orientation of 3c

To establish correlation between the dihedral angle values obtained from X-ray crystallographic data with the results of DFT calculations, the strategy was to calculate the energy of the dimer by adopting gradual change of dihedral angle parameter and to compare those computed energies with X-ray crystallographic results using **Gaussian 09W** where mpwb95/6-31++G was used as basis set. Dihedral angle value at the minimum of the graph obtained from DFT calculated results indicated most stable  $\pi$ - $\pi$  stacking association, corresponding to the optimized dihedral angle in each case of crystal lattice for skew-*syn* and skew-*anti* dispositions exhibited in the molecules **3b** and **3c**. Graphical representation of the correlation between the DFT calculated optimized dihedral angle and X-ray crystallographic results for skew-*syn* arrangement in **3b** and skew-*syn* and skew-*anti* dispositions in **3c** are shown in Fig. 20 and Fig. 21 respectively (total data of DFT calculation are shown in Supplementary data, Table S30-S32).



**Fig. 20.** a) shows dihedral angel for skew-*syn* orientation of **3b** obtained from X-ray crystallographic analysis and b) shows plot of energy of the stacked dimer against varying dihedral angel parameter (keeping slip angel intact).



**Fig. 21.** a), b) show dihedral angel for skew-*syn* orientation and skew-*anti* orientation of **3c** obtained from X-ray crystallographic analysis and b), d) show plot of energy of the stacked dimer against varying dihedral angel parameter (keeping slip angel intact).

In each case, DFT calculated values were in good agreements with the X-ray crystallographic results

(shown in Table 13).

# Table 13

Comparison table of dihedral angel obtained from X-ray crystallographic results and DFT-calculatios for **3b** and **3c** 

	Compound	3b	3c	
Dihedral angel for skew- <i>syn</i>	Dihedral angel obtained from X-ray crystallographic result (°)	29.25	41.27	
orientattion	DFT-calculated optimized dihedral angel (°)	26	41	
Dihedral angel for skew- <i>anti</i> orientattion	Dihedral angel obtained from X-ray crystallographic result (°)	149.54	136.35	
	DFT-calculated optimized dihedral angel (°)	-	137	

#### 3.4. Halogen-halogen interaction and molecular channel formation of 2a-e and 3a-e

Compounds 2d and 2e had shown halogen-halogen soft interactions in their crystal lattices. Although there exist two types of halogen-halogen soft interactions, both 2d and 2e display type-I interaction only. This type-I interaction is characterized by an inter-halogen distance being generally less than the sum of van der Waals radii of two halogens involved and a symmetrical array of R-X<sub>1</sub>...X<sub>2</sub>-R with  $\theta_1 = \theta_2$  (where  $\theta_1$  and  $\theta_2$  are the R-X<sub>1</sub><sup>...</sup>X<sub>2</sub> and X<sub>1</sub><sup>...</sup>X<sub>2</sub>-R angles respectively). Although there exists disagreement about the exact nature of the halogen-halogen interaction, it has to be governed by attractive forces [11,58] or by minimization of repulsive forces [59-62]. DFT studies [63] have indicated that the most stabilizing situations correspond to  $\theta_1 = \theta_2 \approx 150$  and the efficiency of the halogen-halogen soft interaction decreases in the order I<sup>...</sup>I>Br<sup>...</sup>Br>Cl<sup>...</sup>Cl. In the crystal lattices of **2e** the I<sup>...</sup>I distance and the  $\theta_1 = \theta_2$  angles had been found out to be 3.705 Å and 159.5° respectively (Fig. 22a) whereas the corresponding values in case of 2d are 3.937 Å and 138.68° (Fig. 22b). These values indicated clearly that the I<sup>...</sup>I interactions was more effective in case of 2e than that of the Br. Br interaction in case of 2d. Such halogen-halogen interaction had not been observed in the lattices of 2c or 2b where the corresponding halogen, chlorine or fluorine, was more electronegative. This effective halogen-halogen interaction had possibly led to the formation of molecular channels lined with bromines in the lattice of 2d (Fig. 22d) and those lined with iodines and carboxamido moieties (Fig. 22c) in the lattice of 2e.





**Fig. 22.** a) and d) show halogen-halogen distances and C-X<sup>...</sup>X angles for **2e** and **2d** respectively ; c) and d) show molecular channels in the lattices of **2e** and **2d** respectively.

Type-II halogen-halogen soft interaction had been observed in the crystal lattice of **3d**. In the crystal lattice of **3d**, the Br<sup> $\dots$ </sup>Br distance and the  $\theta_1$  and  $\theta_2$  angles were measured as as 3.95 Å, 158.87° and 87.44° respectively (Fig. 23a, Fig. 23b). Such halogen-halogen interactions were not identified in the lattices of **3b**, **3c** and **3e** where the corresponding halogens were fluorine, chlorine and iodine respectively. This effective halogen-halogen interaction has led to the formation of molecular channels in the lattice of **3d** along c-direction (Fig. 23c).



Fig. 23. a) and b) show halogen-halogen distances and C-X<sup>...</sup>X angles for 3d respectively, c) shows dihedral angles of C<sub>1</sub>-Br<sub>1</sub><sup>...</sup>Br<sub>1</sub>-C<sub>1</sub> for 3d, d) shows molecular channels in the lattices of 3d along c-direction.</sup>

Although halogen-halogen soft interaction was absent in case of **3e**, it had shown molecular ordering in its lattice generating arrays of the molecular channels along c-direction (shown in Fig. 24)



Fig. 24. Molecular channel arrangement in the lattices of 3e along c-direction. *3.5. Nanostructural behaviour* 

During the years of new millennium, one dimensional (1-D) organic nano-materials based on small molecules, have attracted great attention, specially for the electronic and optoelectronic devices [64]. 1-D organic nano-structures have been very efficiently constructed with the help of their  $\pi$ - $\pi$  stacking [65] and/or hydrogen bonding [67] properties. Nano-structural materials like tubes [67], rods [68] or belts [69] have been under extensive investigations due to their properties, much different from their bulk versions. Reportedly a 1-D assembly can be obtained very easily by dispersing a concentrated solution of a 'promising' nano-material into a 'poor solvent'[70].

Single crystal X-ray crystallographic studies of series of 5-amino-1-(phenyl/p-halophenyl)imidazole-4carboxamides (**2a-e**) and 5-amino-1-(phenyl/p-halophenyl)imidazole-4-carbonitriles (**3a-e**) had shown different modes of  $\pi$ - $\pi$  stacking interactions, extensive hydrogen bonding network and soft halogenhalogen interactions. These observations indicated to investigate the formation of nano-structural behaviour of those compounds. To explore the influence of solvent polarity on morphology, the morphological heterogeneity experiments of the compounds **2a-e** and **3a-e** using different solvent systems with the change of polarities had been performed. Scanning electron microscopic (SEM) studies revealed that **2a-e** and **3a-e** had given self-assembled nanostructures in non-polar solvent benzene only. The selfassembled morphological heterogeneity had been checked by preparation of the sample through dispersing the compounds (5Mm) by way of sonication for 2 hours at room temperature in a series of solvents such **41** 

as pet ether, methanol, acetonitrile and benzene. The respective solution was allowed to drop on glass cover slips and allowed to dry slowly for 48 hours. Then the SEM images were taken. SEM images revealed that the formation of 1-D nanostructures had been successfully constructed for **2a-c**, and **3a-e** in benzene except **2d** which formed nano-vesicles in this particular solvent benzene. Minimum widths of nano-rod in benzene had been found to be ~92.0nm, 111.57nm, 90nm and 268.70nm for **2a-c** and **2e** respectively (Fig. 25). The SEM images of nano-rod formation are shown in Fig. 25.



**Fig. 25.** a), b), c) and d) show SEM images as nano-rods arising from **2a**, **2b**, **2c** and **2e** in benzene Although, nano-rod morphologies had been observed for **2a-c** and **2e** in benzene, **2d** formed nano-vesicle morphology in benzene having size distribution ~39nm (Fig. 26).



#### Fig. 26. SEM images of 2d in benzene

1-D nanostructure morphology was also viewed for **3a-e** in benzene. Minimum widths of nano-rod in benzene were found to be ~155.86nm, 276.24nm, 70.56nm, 94.67nm and 138.01nm for **3a**, **3b**, **3c**, **3d** and **3e** respectively (shown in Fig. 27).



Fig. 27. a), b), c), d) and e) show SEM images as nano-rods arising from 3a, 3b, 3c, 3d and 3e in benzene

#### 4. Conclusion

In conclusion I had successfully chosen a composite system of imidazole, characteristic of an AICA, with a phenyl/p-halophenyl substituent at N-1 so that the imidazole and the phenyl together constitute a freerotable system. All the four halogens (F,Cl,Br,I) had been considered as the *p*-substituents of phenyl to unravel the profound effect of halogens on the supramolecular architecture in the lattices of the compounds by way of their X-ray crystallographic and DFT studies. In their crystal lattices, phenyl/pfluorophenyl/p-chlorophenyl compounds 2a-c had shown anti-parallel imidazole-imidazole stackformations. On the other hand, p-bromo/p-iodophenyl compounds 2d, 2e avoided such anti-parallel imidazole-imidazole stacking in their crystal lattices. Each of 2a-c and 2e had shown anti-parallel phenylphenyl  $\pi$ - $\pi$  stacking of PD-type; 2d (with bromo as the halogen) indicated *anti*-parallel Sandwich stacking, a choice appearing unusual from the standpoint of stabilization. Compounds **3b-e** had shown T-stacking interactions along with *anti*-parallel PD-type imidazole-imidazole and phenyl-phenyl  $\pi$ - $\pi$  stacking as well as specifically compound 3a revealed dual T-stackings interactions along with anti-parallel PD-type interaction involving imidazole-imidazole stacking. Halogen-halogen soft interactions and formation of molecular channels were found to play an important role in the series of our compounds. In each of the cases of  $\pi$ - $\pi$  anti-parallel and T-stacking arrangements, DFT studies had been taken up for the purpose of quantification of the  $\pi$ - $\pi$  stacking and T-stacking stabilization energy. Detailed DFT calculations of the optimized energies of the dimer had been undertaken by taking out the closest stacked pair from the lattice and gradually changing the stacking distance parameter. When the graph of stacking energy vs. stacking distance was analyzed, minimum of the curve corresponded to the stacking energy of the dimer. In each case, DFT calculated results were in good agreements with X-ray crystallographic results. Similar DFT studies on simultaneous syn-parallel imidazole and phenyl dimer systems corresponded to no  $\pi$ - $\pi$  stacking stabilization at all. Nano-structural behaviour of compounds 2a-e and 3a-e had been observed in non-polar

solvent benzene. Compounds **2a-c**, **2e** and **3a-e** had shown excellent -D nano-structural properties but **2d** had shown nano-vesicle morphology in benzene.

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# Appendix A. Supplementary materials

Supplementary data associated with this article can be found, in the online version, at

(http:////....).

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- Synthesis of 1-(phenyl/p-halophenyl)AICA and its nitrile derivatives.
- Study the change in stacking patterns with functional group change.
- Role of halogens for fixing the stacking patterns.
- DFT study to quantify T-stacking and  $\pi$ - $\pi$  stacking of imi-imi and phe-phe.
- Investigation of nano-structural behavior.