

Recurrent Supramolecular Motifs in a Series of Acid–Base Adducts Based on Pyridine-2,5-Dicarboxylic Acid *N*-Oxide and Organic Bases: Inter- and Intramolecular Hydrogen Bonding

Masoud Mirzaei,* Fereshteh Sadeghi, Krešimir Molčanov, Jan K. Zareba,* Rosa M. Gomila, and Antonio Frontera*



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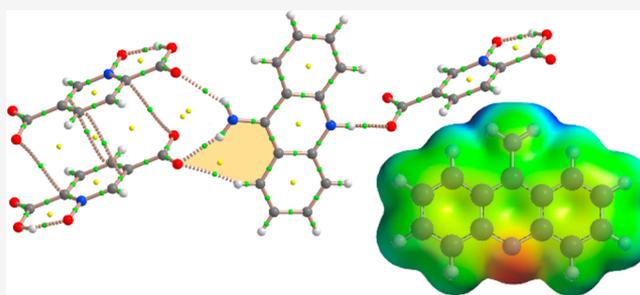


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ABSTRACT: Salts and ionic cocrystals simultaneously comprising *N*-oxide and carboxylic acid functional groups constitute a very fertile ground for the investigation of various proton transfer phenomena. This is because such compounds combine two types of proton transfer: that is, inter- and intramolecular hydrogen bonding in acid–base systems. To this end, a series of novel salts based on pyridine-2,5-dicarboxylic acid *N*-oxide (H₂pydco) as an organic acid and 2,4,6-triamino-1,3,5-triazine (tata), 2-amino-pyrimidine (2a-pym), 2-amino-6-methylpyridine (2a-6mpy), 1,10-phenanthroline (phen), and 9-aminoacridine (9a-acr) as organic bases have been synthesized and characterized by elemental analyses, infrared spectroscopy, and single-crystal X-ray diffraction: (Htata)⁺(Hpydco)[−] (1), (H2a-pym)⁺(Hpydco)[−] (2), (H2a-6mpy)⁺(Hpydco)[−] (3A,B), [(Hphen)⁺(Hpydco)[−]](H₂pydco) (4), and [(H9a-acr)⁺(Hpydco)[−]].EtOH (5). The unit cells of 3A and 3B differ slightly; however, 3A crystallizes in a chiral orthorhombic space group *P*₂₁₂₁, while 3B crystallizes in the achiral space group *P*₂₁/*n*. In both cases, the asymmetric unit comprises one cation and one anion. The influence of different organic cations on the packing of Hpydco[−] in the crystal lattice is studied. The most important feature of these crystals is the presence of extensive O–H⋯O, N–H⋯O, N–H⋯N, and C–H⋯O hydrogen bond networks, which form base-dependent supramolecular synthons: 1, 2, and 3A,B comprise an α -aminopyridinium moiety, and all involve the hydrogen-bonded motif *R*₂²(8) with the Hpydco[−] anion. Compounds 4 and 5, which lack the α -aminopyridinium moiety, reveal different hydrogen-bonding patterns. The interaction energies of each individual hydrogen bond have been estimated using the quantum theory of “atoms-in-molecules”, which led us to the identification of the energetically favorable antielectrostatic N–H⋯N hydrogen bonds (stabilization energy of 4.0 kcal/mol) between positively charged melaminium species in 1. It has been also established that charge-assisted hydrogen bonding does not always offer an energetic advantage over “noncharged” hydrogen bonds. With the use of Hirshfeld surface (HS) analysis we have also explored the influence of the protonation state of pydco species on the composition of contact contributions, as well as established specific properties of their 2D fingerprint plots. Finally, a comment is provided on the applicability of HS analysis for the exploration of polymorphs featuring intramolecular proton transfer.



INTRODUCTION

Crystalline solids are formed when molecules arrange in a very orderly fashion by means of many types of intermolecular interactions such as classical and nonclassical hydrogen bonding, halogen bonding, π stacking, ion pairing, and van der Waals and donor–acceptor interactions, to name only a few. The importance of crystal engineering is highlighted when our purpose is to obtain a compound with desired solid-state properties, to understand the stability (or its lack) of a given crystalline network or structural motif, or to answer the question: “How do crystalline solids self-assemble?”. For this purpose, one of the most leading and efficient methods is the use of single-crystal X-ray diffraction for experimental structure determination and confrontation of the obtained results with the knowledge already garnered by crystal engineers.^{1–4} In

cases where different crystalline structures are obtained but feature the same composition (polymorphism phenomenon), crystallographic structural analysis is one of the most effective research methods that allow one to resolve the fine details of the molecular environment of such species. Generally, polymorphs have different properties such as melting point, stability, density, crystal habit, etc. Polymorphs are classified into several subcategories, such as conformation, packing, synthon, structural polymorphism, etc. Furthermore, poly-

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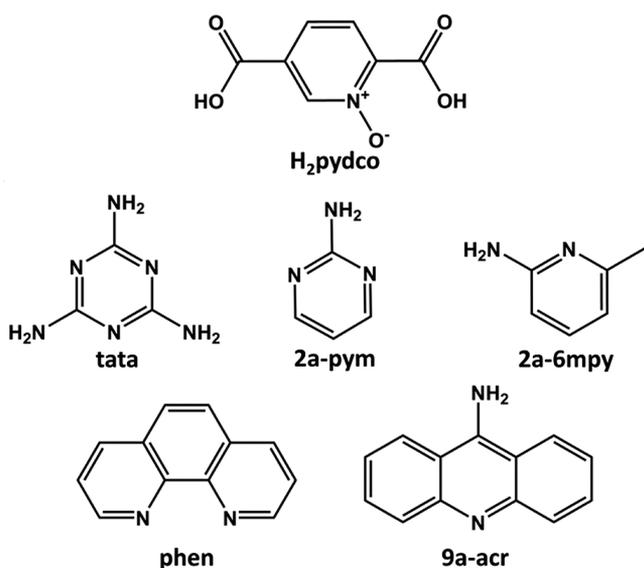
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morphism is not limited to single-component crystal structures only; on the contrary, salts are a very fertile area for the formation of polymorphic structures. Indeed, a salt formation strategy can be used to develop drugs to enhance their physicochemical and clinical properties, such as solubility in water, dissolution rate, chemical stability, tableting, etc. At least half of the marketed active pharmaceutical ingredients (APIs) are administered as salts. In general, a salt is formed via proton transfer from an acid to a base, leading to the formation of a conjugate acid (protonated base) and conjugate base (deprotonated acid).^{5–7} Over the past decade, Mirzaei et al. have focused on the synthesis and structural characterization of novel supramolecular proton-transferred compounds^{8–13} and coordination complexes^{14,15} based on derivatives of pyridine-dicarboxylic acids, owing to their remarkable applications such as anticancer properties, anti-HIV agent, gas adsorbent, etc.^{16–23}

In this paper, we report on the synthesis and X-ray characterization of a series of new salts with N-containing organic aromatic bases such as 2,4,6-triamino-1,3,5-triazine (tata), 2-aminopyrimidine (2a-pym), 2,6-diaminopyridine (2a-6mpy), phenanthroline (phen), and 9-aminoacridine (9a-acr) (Scheme 1) and pyridine-2,5-dicarboxylic acid *N*-oxide

Scheme 1. (Top) Structural Formula of H₂pydco and (Bottom) Structural Formulas and Abbreviations Used for Aromatic Bases



(H₂pydco) as the organic acid, using a proton transfer mechanism: (Htata)⁺(Hpydco)[−] (**1**), (H2a-pym)⁺(Hpydco)[−] (**2**), (H2a-6mpy)⁺(Hpydco)[−] (two polymorphs, **3A** and **3B**), [(Hphen)⁺(Hpydco)[−]](H₂pydco) (**4**), and [(H9a-acr)⁺(Hpydco)[−]].EtOH (**5**). It is generally accepted that, if there is an adequate pK_a difference between the COOH of the aromatic acid and the nitrogen atom of the aromatic amine, a proton transfer phenomenon will occur and will likely generate the ionic hydrogen bond N⁺–H⋯O[−].^{22,23} Single crystals of salts **2–5** were obtained in the presence of Mn(II) ion, but no coordination compounds have been isolated, probably due to the preferential formation of stable, extended networks of hydrogen bonds, supported by the process of proton transfer.^{24,25} In that context it is worth adding that only four complexes with H₂pydco and lanthanoids were found in the

Cambridge Structural Database (CSD version 5.40 updates (November 2018)).²⁶ Given the tendency of H₂pydco to participate in complex hydrogen-bonded networks, in this contribution, we decided to more closely explore the aforementioned acid–base adducts (five salts and one ionic cocrystal) involving H₂pydco and characterize their supramolecular behavior. We place a particular focus on the determination of recurrent supramolecular motifs that emerge from their intermolecular and intramolecular hydrogen bonding patterns. Insight into these properties is obtained with the use of classical structure description involving determination of hydrogen bond ring and chain motifs as well as Hirshfeld surface analysis. Experimental observations are corroborated by a DFT study, which offers the opportunity for evaluation of the energetic contribution of hydrogen bonds, which are formed between the counterions and also between the units of the same sign (cation⋯cation and anion⋯anion).

METHODS

General Methods and Materials. All chemicals and solvents used for the syntheses were of reagent grade and were purchased from Merck Chemicals and used without further purification. The organic acid (H₂pydco) was synthesized according to a reported procedure.¹⁵ Melting points were determined using a Barnstead Electrothermal 9300 apparatus. IR spectra were recorded from KBr pellets in the 4000–400 cm^{−1} region using a Buck 500 IR spectrometer. Elemental analysis (CHN) was performed using a Thermo Finnigan Flash-1112EA microanalyzer.

Crystallographic Analyses. Single-crystal measurements were performed on an Oxford Diffraction Xcalibur Nova R (microfocus Cu tube) equipped with an Oxford Instruments CryoJet liquid nitrogen cooling device. The program package CrysAlis PRO (Agilent, 2011)²⁷ was used for data reduction and numerical absorption correction. The structures were solved using SHELXS97²⁸ and refined with SHELXL-2017.²⁸ The models were refined using full-matrix least-squares refinement; all non-hydrogen atoms were refined anisotropically. Hydrogen atoms were located in a difference Fourier map and refined as either free entities or a mixture of free and riding entities. Water molecules were refined using the following restraints: O–H bonds were restrained to 0.95(2) Å, and H⋯H distances were restrained to 1.50(4) Å.

Synthesis of (Htata)⁺(Hpydco)[−] (1**).** A solution of H₂pydco (0.037 g, 0.20 mmol) in ethanol/water (1/1; 10 mL) was added dropwise to a solution of tata (0.025 g, 0.20 mmol) in deionized water (5 mL) and stirred for 4 h. After 1 week, colorless block-shaped crystals of **1** were obtained in about 47% yield (based on H₂pydco) by slow evaporation. (mp >300 °C). Anal. Calcd for C₁₀H₁₁N₇O₅: C, 38.84; H, 3.59; N, 31.71. Found: C, 39.81; H, 3.38; N, 32.05. IR (KBr pellet, cm^{−1}): 3427(m), 3297(m), 1674(s), 1645(s), 1512(m), 1372(s), 1228(w).

Synthesis of (H2a-pym)⁺(Hpydco)[−] (2**).** A solution of H₂pydco (0.029 g, 0.16 mmol), MnO₂ (0.007 g, 0.08 mmol), and 2a-pym (0.015 g, 0.16 mmol) in ethanol/water (1/1; 25 mL) was prepared and stirred for 4 h under mild conditions. After 10 days colorless prism-shaped diffraction-quality single crystals of **2** were obtained and collected through a slow evaporation of the solvent under ambient conditions in about 57% yield (based on H₂pydco) (mp 205 °C). Anal. Calcd for C₁₁H₁₀N₄O₅: C, 47.49; H, 3.62; N, 20.14. Found: C, 48.73; H, 3.28; N, 21.38. IR (KBr pellet, cm^{−1}): 3292(br), 2704(w), 1972(m), 1707(s), 1679(s), 1552(m), 1370(s), 1263(m), 1228(w).

Synthesis of (H2a-6mpy)⁺(Hpydco)[−] (3A**).** A solution of H₂pydco (0.029 g, 0.16 mmol), MnCl₂·2H₂O (0.013 g, 0.08 mmol), and 2a-6mpy (0.017 g, 0.16 mmol) in ethanol/water (3/1; 40 mL) was prepared and stirred for 4 h under mild conditions. After 15 days yellow prism-shaped diffraction-quality single crystals of **3A** were obtained and collected through a slow evaporation of the solvent under ambient conditions in about 45% yield (based on H₂pydco) (mp 185 °C). Anal. Calcd for C₁₃H₁₃N₃O₅: C, 53.61; H, 4.50; N,

Table 1. Crystal Data and Data Collection and Refinement Parameters for 1–5

	1	2	3A	3B	4	5
Crystal Data and Data Collection Parameters						
empirical formula	C ₁₀ H ₁₁ N ₇ O ₅	C ₁₁ H ₁₀ N ₄ O ₅	C ₁₃ H ₁₃ N ₃ O ₅	C ₁₃ H ₁₃ N ₃ O ₅	C ₂₆ H ₁₈ N ₄ O ₁₀	C ₂₂ H ₂₁ N ₃ O ₆
formula wt	309.26	278.23	1165.06	291.26	546.44	399.36
cryst syst	triclinic	monoclinic	orthorhombic	monoclinic	triclinic	triclinic
space group	$\bar{P}1$	$P2_1/n$	$P2_12_12_1$	$P2_1/n$	$\bar{P}1$	$\bar{P}1$
<i>a</i> (Å)	8.2019(4)	3.7453(1)	6.6681(1)	7.9463(2)	6.9340(2)	7.2141(6)
<i>b</i> (Å)	8.7490(6)	10.8274(3)	7.9560(1)	6.6606(1)	7.2620(3)	11.1551(6)
<i>c</i> (Å)	10.0972(6)	28.1248(8)	25.7764(4)	25.8445(5)	23.590(8)	13.2745(10)
α (deg)	95.149(5)	90	90	90	85.019(3)	69.808(7)
β (deg)	109.230(5)	90.814(3)	90	93.909(2)	87.751(3)	89.600(7)
γ (deg)	114.764(6)	90	90	90	75.820(3)	75.762(7)
<i>V</i> (Å ³)	598.91(6)	1140.40(5)	1367.47(3)	1364.69(5)	1147.17(7)	968.23(12)
<i>Z</i>	2	4	1	4	2	2
μ (mm ⁻¹)	1.22	1.13	0.94	0.94	1.06	0.87
no. of measd, indep, and obsd indices (<i>I</i> > 2 σ (<i>I</i>)) rflns	6407, 2262, 1849	5014, 2123, 1922	6662, 2721, 2587	11674, 2807, 2584	10319, 4698, 4195	8494, 3958, 3210
<i>R</i> _{int}	0.046	0.022	0.026	0.036	0.029	0.022
(sin θ/λ) _{max} (Å ⁻¹)	0.618	0.628	0.628	0.629	0.629	0.630
Refinement Parameters						
<i>R</i> (<i>F</i> ² > 2 σ (<i>F</i> ²)), <i>R</i> _w (<i>F</i> ²), <i>S</i>	0.051, 0.160, 1.04	0.051, 0.155, 1.11	0.038, 0.109, 1.04	0.046, 0.141, 1.09	0.045, 0.134, 1.05	0.073, 0.246, 1.04
no. of rflns	2262	2123	2721	2807	4698	3958
no. of params	207	190	190	194	362	286
no. of restraints	0	0	0	0	0	30
$\Delta\rho_{\max}$ $\Delta\rho_{\min}$ (e Å ⁻³)	0.31, -0.42	0.24, -0.16	0.17, -0.18	0.21, -0.19	0.29, -0.19	0.73, -0.28

14.43. Found: C, 54.47; H, 4.23; N, 14.65. IR (KBr pellet, cm⁻¹): 3292(m), 2798(w), 1981(m), 1695(m), 1498(w), 1365(s), 1249(m).

Synthesis of (H2a-6mpy)⁺(Hpydco)⁻ (3B). A solution of H₂pydco (0.029 g, 0.16 mmol), MnCl₂·2H₂O (0.026 g, 0.16 mmol), and 2a-6mpy (0.034 g, 0.32 mmol) in ethanol (10 mL) was prepared and stirred for 4 h under mild conditions. After 10 days colorless prism-shaped diffraction-quality single crystals of **3B** were obtained and collected through a slow evaporation of the solvent under ambient conditions in about 60% yield (based on H₂pydco) (mp 199 °C). Anal. Calcd for C₁₃H₁₃N₃O₅: C, 53.61; H, 4.50; N, 14.43. Found: C, 54.51; H, 4.12; N, 14.60. IR (KBr pellet, cm⁻¹): 3283(br), 1971(m), 1693(s), 1635(m), 1585(m), 1379(s), 1252(m), 1223(w).

Synthesis of [(Hphen)⁺(Hpydco)⁻](H₂pydco) (4). A solution of H₂pydco (0.037 g, 0.20 mmol), MnCl₂·2H₂O (0.032 g, 0.20 mmol), and 1,10-phenanthroline hydrate (0.040 g, 0.20 mmol) in ethanol/water (1/1; 25 mL) was prepared and stirred for 4 h under mild conditions. After 20 days colorless prism-shaped diffraction-quality single crystals of **5** were obtained and collected through a slow evaporation of the solvent under ambient conditions in about 47% yield (based on H₂pydco) (mp 200 °C). Anal. Calcd for C₂₆H₁₈N₄O₁₀: C, 57.15; H, 3.32; N, 10.25. Found: C, 57.33; H, 3.15; N, 10.38. IR (KBr pellet, cm⁻¹): 3446(br), 3076(m), 1678(s), 1537(m), 1393(s), 1245(m).

2.8. Synthesis of [(H9a-acr)⁺(Hpydco)⁻](EtOH) (5). A solution of H₂pydco (0.37 g, 0.20 mmol), MnCl₂·2H₂O (0.016 g, 0.10 mmol), and 9a-acr (0.019 g, 0.10 mmol) in ethanol/water (1/1; 30 mL) was prepared and stirred for 8 h under mild conditions. Yellow needle-shaped crystals were obtained after 2 weeks in about 61% yield (based on H₂pydco) by slow evaporation (mp 244 °C). Anal. Calcd for C₂₂H₂₁N₃O₆: C, 62.41; H, 5.00; N, 9.92. Found: C, 63.05; H, 4.70; N, 10.86. IR (KBr pellet, cm⁻¹): 3456(br), 3172(m), 1742(m), 1655(s), 1615(s), 1548(w), 1407(m), 1263(m), 1225(w).

Theoretical Methods. All DFT calculations were carried out using the Gaussian-16 program²⁹ at the B3LYP-D3/def2-TZVP level of theory and using the crystallographic coordinates (only the positions of the H atoms have been optimized). Atoms in molecules (AIM)³⁰ analysis was performed at the same level of theory. The calculation of AIM properties was done using the AIMAll program.³¹

RESULTS AND DISCUSSION

Infrared Spectroscopy. The IR spectra of H₂pydco and all studied compounds are shown in Figure S1 in the Supporting Information. The absorptions at 3000–3500 cm⁻¹ can be assigned to the NH and OH stretching vibrations of the amino groups or O–H in the carboxylate group of the H₂pydco ligand and water molecules, respectively.^{32,33} The strong band $\nu_{\text{as}}(\text{COO}^-)$ and $\nu_{\text{s}}(\text{COO}^-)$ stretching modes appeared for free H₂pydco at 1726 and 1419 cm⁻¹, respectively, which were shifted in all of the compounds to lower wavenumbers in the ranges 1695–1626 and 1407–1365 cm⁻¹, respectively. Furthermore, in the infrared spectrum of H₂pydco (Figure S1) a strong band at 1230 cm⁻¹ can be attributed to the presence of an N–O group.²⁶ Bands in the 1228–1207 cm⁻¹ region for all compounds were assigned to the stretching vibration of the N–O group.

Description of the Crystal Structures. The crystallographic and collection data for 1–5 are gathered in Table 1. Additionally, selected bond lengths, valence angles, and hydrogen bond geometries are given in Tables S1 and S2 in the Supporting Information. In all studied salts the Hpydco⁻ anion forms an intramolecular hydrogen bond between a carboxyl moiety and the N-oxide group, which can be represented by an S₁¹(6) graph set^{34,35} (Figure 1a). The hydrogen bond is stabilized by resonance (RAHB) of the carboxyl and N–O groups.

Compounds **1**, **2**, and **3A,B** contain an α -aminopyridinium moiety, and all form the hydrogen-bonded motif R₂²(8) with the Hpydco⁻ anion (Figure 1b). An attraction of proton donors and acceptors is augmented here by the negative charge of the carboxylate group of the Hpydco⁻ anion; thus, a pair of resonance- and charge-assisted hydrogen bonds (CAHB, RAHB) is formed. This fragment is a well-known supra-molecular synthon.³⁵ 3D packings of compounds **1**, **2**, **3A,B** are

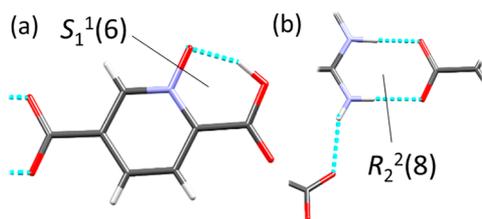


Figure 1. Highly conserved hydrogen bonded motifs: (a) an $S_1^1(6)$ ring present in all Hpydco[−] anions and (b) an $R_2^2(8)$ ring present in **1**, **2**, and **3A,B**.

completed by other proton donors and acceptors forming secondary hydrogen-bonded motifs. Compounds **4** and **5**, which lack the α -aminopyridinium moiety, form different hydrogen-bonding patterns.

Crystal Structure of (Htata)⁺(Hpydco)[−] (1). The asymmetric unit **1** consists of one cation (Htata)⁺ and one anion (Hpydco)[−] (see Figure 2). The intramolecular hydrogen bond in the $S_1^1(6)$ ring is particularly strong (the H2...O3 distance is 1.29(4) Å). The cation and anion are bonded into one-dimensional (1D) infinite chains through five distinct N–H...O intermolecular hydrogen bonds. Within these chains one can identify four supramolecular synthons³⁵ of $R_2^2(8)$, $R_2^2(6)$, $R_1^1(4)$, $R_1^1(4)$, graph set notations (Figure 3). The latter two motifs are components of larger $R_2^2(6)$ graph set. It is obvious that O2 and H6B atoms act as a double acceptor and a bifurcated donor, respectively.

Neighboring chains are linked to each other by N–H...O and N–H...N hydrogen bonds between the anions and cations, generating two-dimensional (2D) HB sheets. We recognize several cyclic synthons in this layer described by $R_4^4(8)$, $R_2^2(8)$, $R_3^3(8)$, and $R_4^4(26)$ graph sets (Figure S2 in the Supporting Information). The three-dimensional (3D) network is built by several weak, but distinct, interactions: (i) π interactions between anionic parts (Hpydco[−]) (3.484 Å), (ii) π interactions between cationic parts (Htata⁺) (3.992 Å), (iii) C–O... π interactions between oxygen of the 5-carboxylate group from Hpydco[−] and the triazine ring of Htata⁺ (3.112 Å), and (iv) N–H... π interactions between nitrogen of the amino group from Htata⁺ and the pyridine ring of Hpydco[−] (3.627 Å) (Figure S3 in the Supporting Information).^{11,36,37}

Crystal Structure of (H2a-pym)⁺(Hpydco)[−] (2). The asymmetric unit of **2** comprises one cation (H2a-pym)⁺ and one anion (Hpydco)[−] (Figure 2). Each anion and cation forms a $R_2^2(8)$ dimer via strong N–H...O hydrogen bonds, and then a 1D linear chain is created by N–H...O (from the amino group, linking to another dimer) and C–H...O hydrogen-bonding interactions along the *b* axis (Figure 4).

A 2D hydrogen-bonded network is created by connection of the linear chains along the *b* axis through N–H...O, C–H...O, and C–H...N hydrogen-bonding interactions along the side view (Figure 5). On the other hand, a linear chain along the *b* axis creates another 2D supramolecular layer via π stacking (with a centroid...centroid separation of 3.745 Å) between pyridyl rings of anion fragments and π stacking (with a centroid...centroid separation of 3.745 Å) between pyrimidyl rings of cation fragments along the *ab* plane (Figure S4 in the Supporting Information).

Crystal Structures of Polymorphs (H2a-6mpy)⁺(Hpydco)[−] (3A,B). Two polymorphs of (H2a-6mpy)⁺(Hpydco)[−] were obtained, through modification of the solvent system combined with a change in the ratio of the

reactants. Their unit cell parameters differ slightly (Table 1); however, **3A** crystallizes in the chiral orthorhombic space group $P2_12_12_1$, while **3B** crystallizes in the achiral space group $P2_1/n$. In both cases the asymmetric unit comprises one cation and one anion.

In both crystal packings structural dimeric motifs of the $R_2^2(8)$ notation are present. In **3A** they are connected by the additional hydrogen bond N–H...O into 1D linear chains along the *a* direction (Figure 6). These linear chains are connected to each other by π stacking (with a centroid...centroid separation of 3.796 Å) between pyridyl rings of anion and cation fragments and thus form 2D layers (Figure 7). Along the *c* axis a 1D ladder is created by N–H...O hydrogen bonds and C–H...O (H4...O3 2.58 Å and H9...O2 2.54 Å) interactions between the anion and cation in the crystal lattice (Figure 8). In **3B** there are additional motifs formed by N–H...O hydrogen bonds: namely, $C_2^1(4)$ and $C_2^2(8)$ (Figure 9). In **3B**, anionic (Hpydco)[−], and cationic (H2a-6mpy)⁺ components of the structure are interdigitating and hold together through (H10...O2; 2.50 Å) interaction. As a result, 1D H-bonded chains are formed (Figure S5 in the Supporting Information). Finally, the linear chains are connected to each other by π stacking (with a centroid...centroid separation of 3.796 Å) between pyridyl rings of anion and cation segments and thus form 2D layers (Figure S6 in the Supporting Information).

Crystal Structure of [(Hphen)⁺(Hpydco)[−]](H₂pydco) (4). Compound **4** is an ionic cocrystal whose asymmetric unit includes one (Hphen)⁺ cation, one (Hpydco)[−] anion, and one neutral H₂pydco (Figure 10). In the main features of the crystal packing one can include hydrogen-bonded double chains parallel to the *b* axis (Figure 10). These chains consist of alternating cationic and anionic components through (H2D...O4A 1.95 Å) hydrogen bonding between the cation (N–H group) and anion (carboxylate group). There are also π interactions (with a centroid...centroid separation of 3.507 Å) between the pyridyl ring of the anion and the phenyl ring of the cation.

Another column is created by neutral H₂pydco, which is linked to the former via O–H...O and C–H...O hydrogen bonds between two anionic and neutral components (Figure 10). The structure of salt **4** is repeated through anionic, neutral, and cationic portions and constructs a 1D linear chain along the side view via strong O–H...O interactions between H₂pydco and Hpydco[−] and C–H...O interactions of the pyridyl rings of Hphen⁺ and oxygen atoms of H₂pydco and Hpydco[−] (Figure S7 in the Supporting Information). The linear chains (Figures S7 and S8 in the Supporting Information) are linked into 2D layers through of C–O... π (a C7AO3A...centroid separation of 3.654 Å) and π interactions (with four different types of centroid...centroid separations of 3.617, 3.661, 3.507, and 3.947 Å) between two rings of Hphen⁺ and Hpydco[−] and N–O... π (an N1O5...centroid separation of 3.766 Å)³⁸ and π interactions (with centroid...centroid separations of 3.794 Å) between pyridyl rings of the H₂phen neutral molecule along the *a* axis.

Crystal Structure of [(H9a-acr)⁺(Hpydco)[−]]-EtOH (5). Single-crystal X-ray diffraction analysis reveals that **5** crystallizes in space group $P\bar{1}$ of the triclinic crystal system. The asymmetric unit contains one (H9a-acr)⁺ cation, one (Hpydco)[−] anion, and one ethanol molecule (Figure 2). Two parallel 1D chains are generated with alternating cationic and anionic components through various N–H...O hydrogen

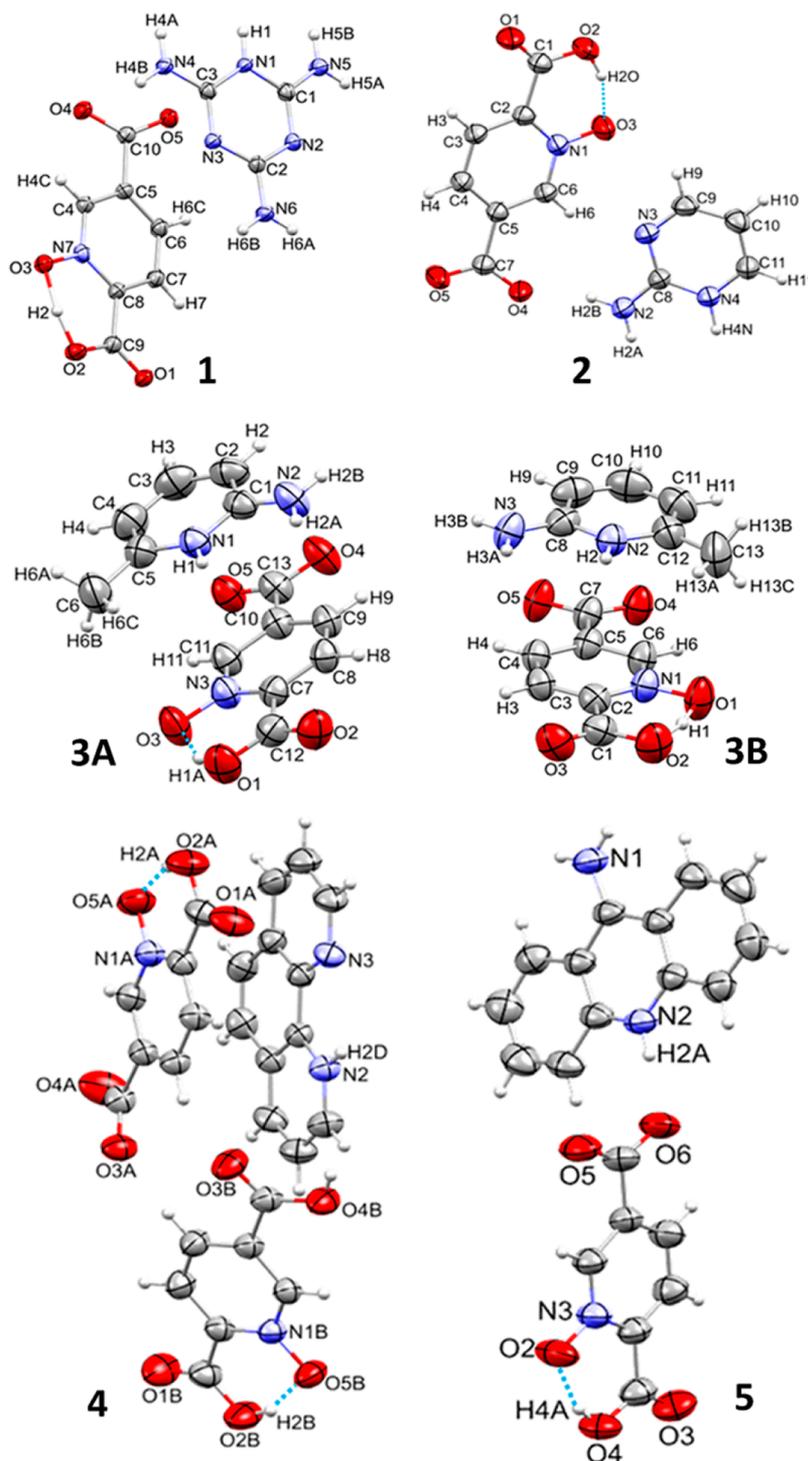


Figure 2. Molecular structures of compounds 1–5 with the displacement ellipsoids drawn at the 50% probability level.

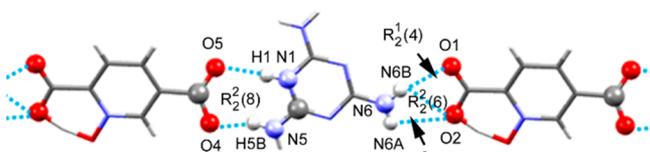


Figure 3. Schematic representation of cyclic motifs created by N–H...O hydrogen bonding in compound 1.

bonds between cations (NH and NH₂ groups) and anions (COO[−] and COOH groups) as well as C–H...O (H2...O4 2.50 Å and H5...O5 2.54 Å) interactions. It should be noted that all the aromatic rings from the (H9a-acr)⁺ cation and (Hpydco)[−] anion feature an antiparallel orientation (see Figure 11).

Adjacent acridinium rings are linked together via π interactions (with two different types of centroid...centroid separations of 3.660 and 3.710 Å) in an AB arrangement, and the pyridyl ring of (Hpydco)[−] participates in π -stacking (with centroid...centroid separation of 3.680 Å) interactions. More-

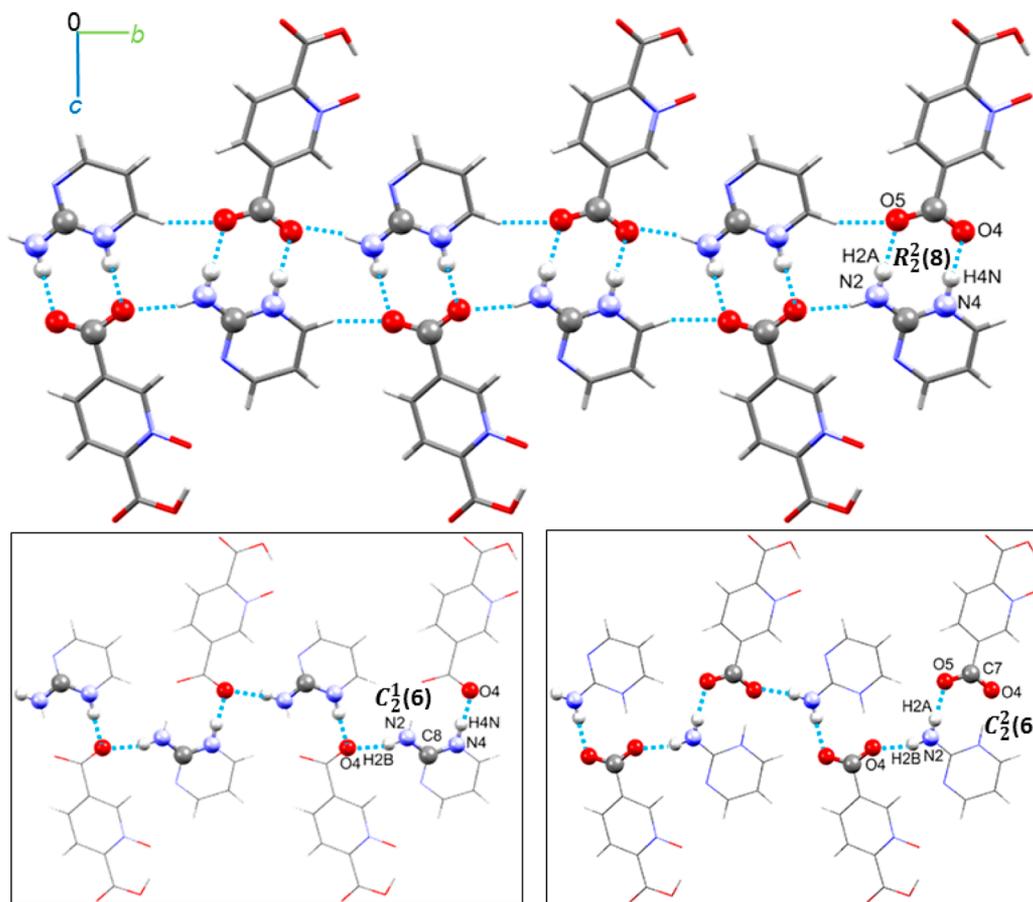


Figure 4. Schematic representation of the 1D linear chain along the *b* axis by N–H...O and C–H...O hydrogen bonding and some graph set notations of **2**.

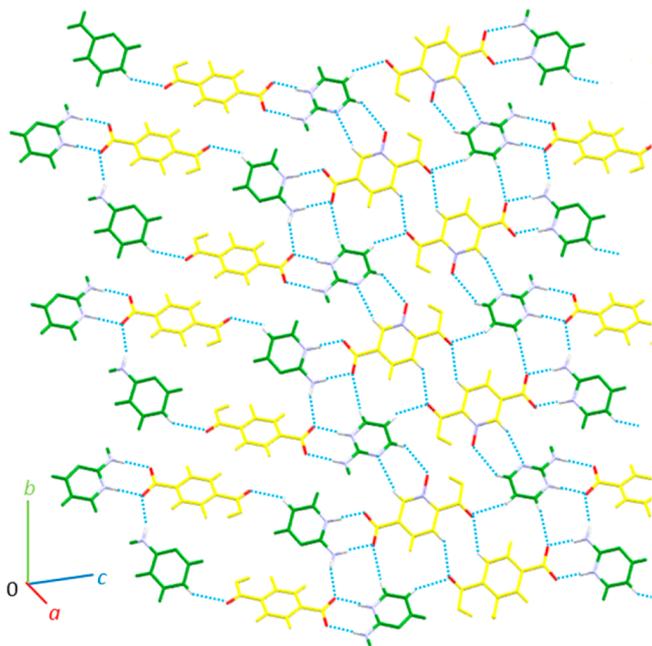


Figure 5. View of the 2D hydrogen-bonded coordination network formed by various hydrogen bonds in **2** along the side view. Symmetry-independent ions are shown in different colors (green and yellow).

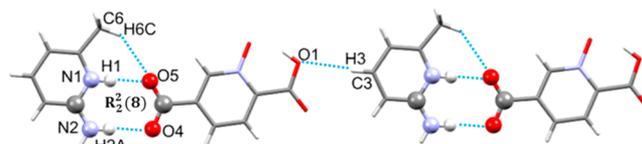


Figure 6. Perspective view of the 1D linear chain and cyclic graph-set notation in **3A** along the *a* direction.

over, in these chains N–H...O hydrogen bonding creates the motifs $C_2^2(8)$, $R_4^4(20)$, and $R_4^4(22)$ (Figure 11). Additionally, these double chains are connected into 2D layers through three types of C–H...O (H9...O2 2.54 Å, H15...O5 2.45 Å, and H18...O3 2.57 Å) interactions and π interactions (with two different types of centroid...centroid separations of 3.799 and 3.915 Å) between all rings of the acridinium cation so that they interact with each other via an ABA arrangement, forming columns along the *a* direction (Figure 12).^{39,40}

Theoretical Study of Noncovalent Interactions. In order to confirm the existence and measure the strength of the H-bonding contacts described above, we carried out DFT calculations and carried out topological analysis of the electron density distribution within the framework of the quantum theory of atoms in molecules (QTAIM method). We have analyzed the supramolecular assemblies of compounds **1–5** and evaluated the contribution of each contact. We and others have successfully used this approach to analyze several types of noncovalent interactions.⁴¹ We have defined energies for the

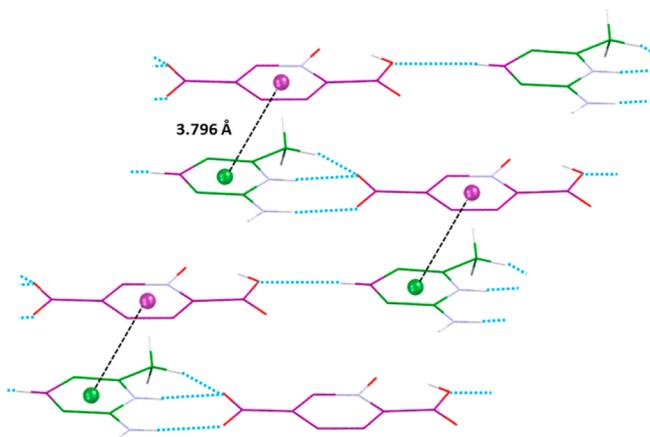


Figure 7. Representation of the 2D layer along the side view of **3A** formed via N–H···O and C–H···O hydrogen-bonding and π interactions. Symmetry-independent molecules are shown in different colors in wireframe style (green and purple), and H atoms bonded to C atoms have been omitted for clarity. The hydrogen bonds and π interactions are shown as blue dotted and black dashed lines, respectively.

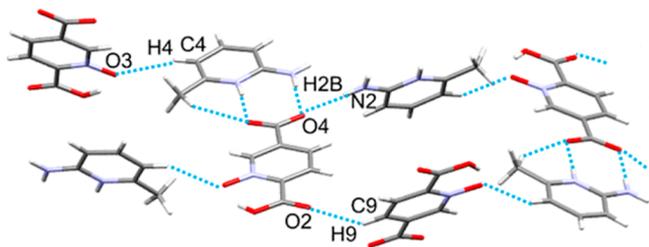


Figure 8. 1D ladder of **3A** viewed along the c axis.

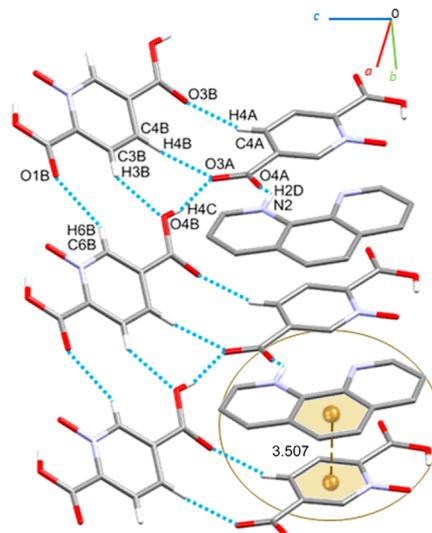


Figure 10. Representation of the 1D chain in **4** formed via O–H···O and C–H···O hydrogen-bonding and π interactions, along the b axis. The hydrogen bonds and π interactions are shown as blue and brown dotted lines, respectively. H atoms bonded to C atoms have been omitted for clarity.

studied contacts according to the conventional approach suggested by Espinosa et al.⁴² and Vener et al.,⁴³ which were developed specifically for HBs.

In **1** different types of N–H···O, N–H···N, and C–H···N intermolecular hydrogen bonds are formed via the three faces of the melaminium cation, as shown in Figure 13. A salt-bridge interaction is formed via the protonated face, where two N–H···O interactions are established. The distribution of critical points and bond paths indicate that each H bond is characterized by a bond critical point (CP) and bond path (BP) that connects the H atom to the N or O atom. In Figure

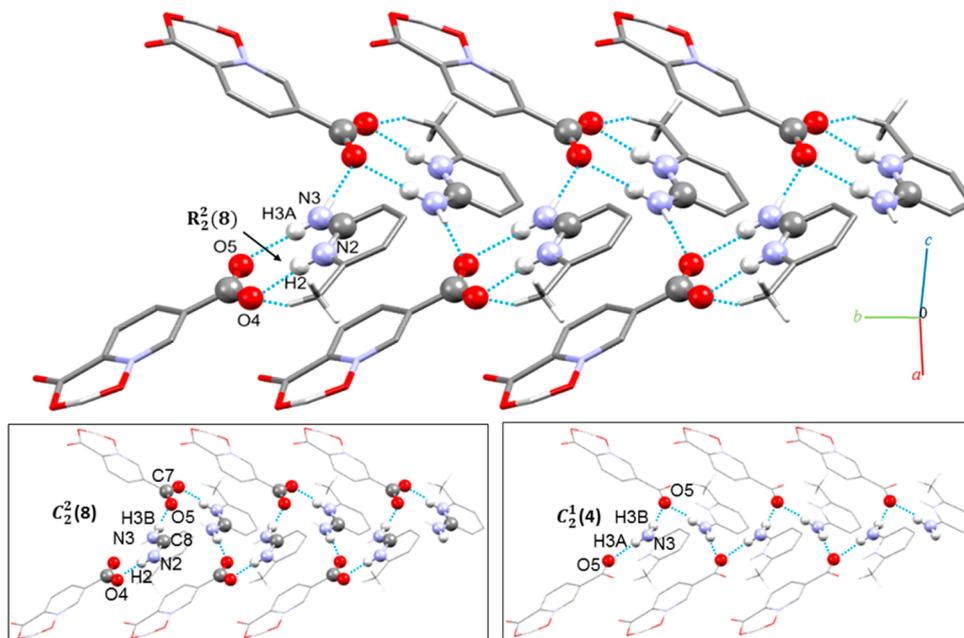


Figure 9. Schematic representation of the 1D zigzag chain along the b axis by N–H···O hydrogen bonding, with indicated graph-set notations $R_2^2(8)$, $C_2^2(8)$, and $C_2^1(4)$ in **3B**. H atoms bonded to C atoms have been omitted for clarity.

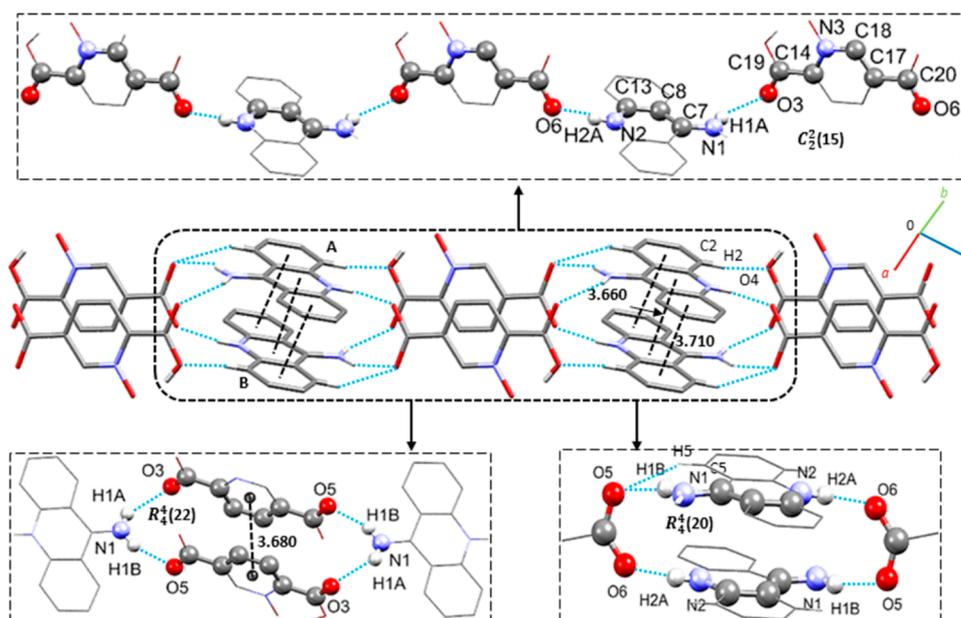


Figure 11. Representation of the 1D double chains along the side view by the N–H···O hydrogen bonding and hydrogen-bonded motifs $C_2^2(8)$, $R_4^4(20)$, and $R_4^4(22)$ in **5**. Distances are shown in Å. The hydrogen bonds and π interactions are shown as blue dotted lines and black dashed lines, respectively. H atoms bonded to C atoms have been omitted for clarity.

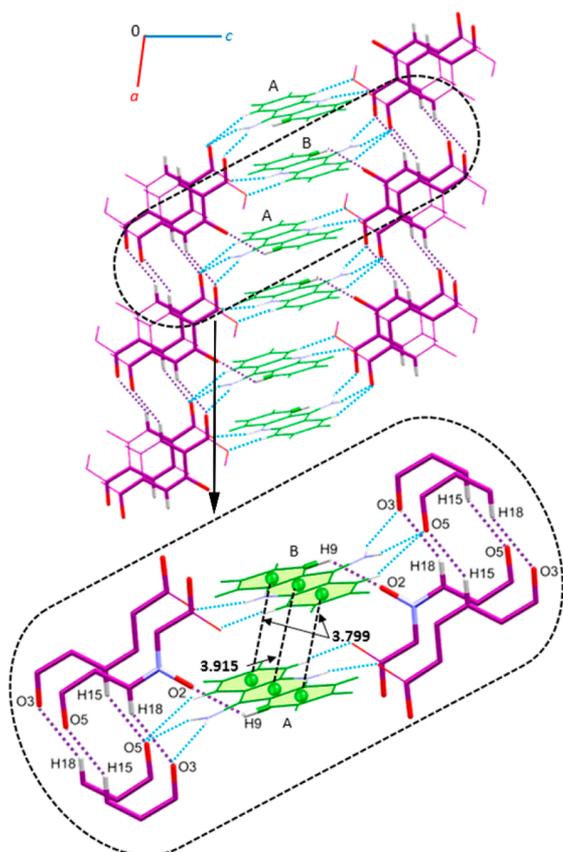


Figure 12. Extended network of the intermolecular interactions of **5**, projected along the *b* direction. Distances are shown in Å. The N–H···O and C–H···O hydrogen bonds and π interactions are shown as blue and purple dotted lines and black dashed lines, respectively. Symmetry-independent ions and are shown in different colors (green, purple).

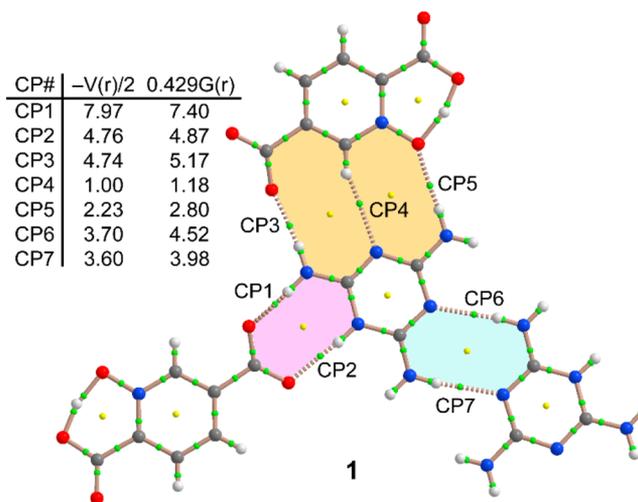


Figure 13. AIM distribution of bond and ring critical points (green and yellow spheres, respectively) and bond paths obtained for the H-bonded assembly of compound **1**. The dissociation energies of the H bonds using the $V(r)$ and $G(r)$ values at the bond CP are indicated (in kcal/mol) in the upper left corner.

13 we also summarize the binding energies (in kcal/mol) for each H bond using both methodologies: i.e., derived from the potential energy density $V(r)$ and from the Lagrangian kinetic energy $G(r)$. The values obtained using both methodologies are similar, and they reveal that the N–H···O H bonds are stronger, apart from that involving the N-oxide (CP5). Interestingly, the antielectrostatic N–H···N H bonds in **1** (see green area in Figure 13) that are formed between the cationic melamine units and characterized by bond critical points CP6 and CP7 (H bonds) are energetically favorable (around 4.0 kcal/mol). Finally, the C–H···N H bond is weak (~ 1.0 kcal/mol), as expected by taking into consideration the H-bond donor group (aromatic C–H bond).

In **2**, we have studied the salt–bridge interaction and also two symmetrically equivalent C–H...O H bonds (characterized by CP3) that connect two (Hpydco)[−] anions (Figure 14).

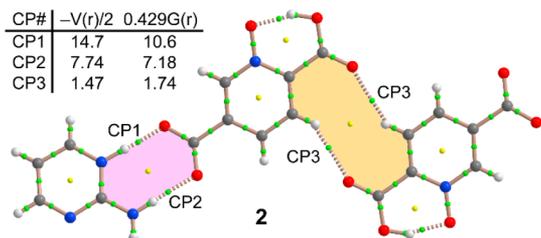


Figure 14. AIM distribution of bond and ring critical points (green and yellow spheres, respectively) and bond paths obtained for the H-bonded assembly of compound **2**. The dissociation energies of the H bonds obtained using the $V(r)$ and $G(r)$ values at the bond CP are indicated (in kcal/mol) in the upper left corner.

The interaction energy of the salt bridge (characterized by CP1 and CP2) is significantly stronger than that observed in **1** (melamine cation), which is likely due to the smallest delocalization of the positive charge in the 2-aminopyrimidinium cation in comparison to melaminium. In fact, the N⁺–H...O bond is stronger than the exocyclic N–H...O bond in **2** and the opposite is observed in **1**, thus confirming the stronger delocalization in the latter. Finally, each antielectrostatic C–H...O H bond established between the anionic moieties contributes 1.47/1.74 kcal/mol.

The QTAIM analyses of polymorphs **3A,B** are shown in Figure 15. In both compounds the strongest H-bonding interaction corresponds to the N⁺–H...O contact of the salt–

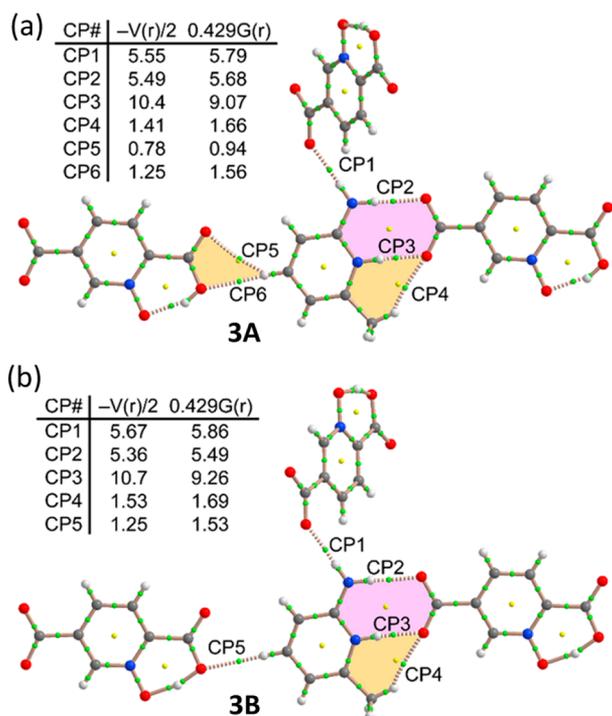


Figure 15. AIM distribution of bond and ring critical points (green and yellow spheres, respectively) and bond paths obtained for the H-bonded assemblies of (a) **3A** and (b) **3B**. The dissociation energies of the H bonds calculated using the $V(r)$ and $G(r)$ values at the bond CP are indicated (in kcal/mol) in the upper left corners.

bridge interaction (CP3), in a manner similar to that for **2**. The presence of a methyl group in the aminopyridine ring assists the salt bridge formation with an additional C–H...O interaction (1.41/1.66 kcal/mol, CP4). The main difference between both polymorphs is the interaction of the cationic moiety with the carboxylic group engaged in the intramolecular H-bond with the N-oxide. In **3A**, the QTAIM analysis suggests the formation of a bifurcated H bond with the C–H in a para position relative to the protonated N atom, whereas in compound **3B** the interaction is characterized by the presence of a single bond CP and BP. In any case, these C–H...O contacts are the weakest interactions of both assemblies, which is in line with previous results.

In **4** the salt–bridge interaction cannot be formed, in contrast to **1**, **2**, and **3A,B**, where it is the most favored interaction. Instead, the anionic carboxylate group interacts with both the neutral H₂pydco moiety and the monoprotonated 1,10-phenanthroline cation (see Figure 16). Un-

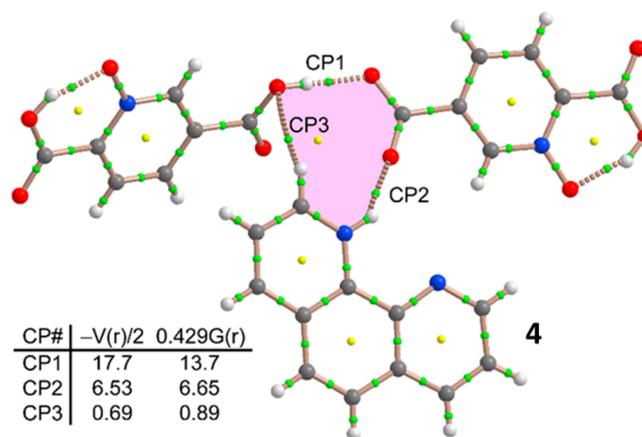


Figure 16. AIM distribution of bond and ring critical points (green and yellow spheres, respectively) and bond paths obtained for the H-bonded assembly of compound **4**. The dissociation energies of the H bonds obtained using the $V(r)$ and $G(r)$ values at the bond CP are indicated (in kcal/mol) in the upper left corner.

expectedly, the strongest interaction corresponds to the O–H...O H bond (CP1) instead to the electrostatically enhanced (CP2) N⁺–H...O H bond. Finally, the distribution of bond CPs and BPs also reveals the existence of an additional and weak C–H...O interaction (CP3, 0.69/0.89 kcal/mol) involving an aromatic C–H group of the 1,10-phenanthroline cation and the carboxylic group.

Figure 17 displays the H-bonded interactions involving the protonated 9-aminoacridine. The amino group establishes two H bonds with two (Hpydco)[−] units (characterized by CP1 and CP2 critical points). The H bond characterized by the critical point CP2 is stronger than that of CP1 due to the anionic nature of the H-bond acceptor in the former (carboxylate). The (Hpydco)[−] units are also stabilized by the formation of π stackings which are characterized by four bond CPs and bond paths interconnecting both anions. As expected, the strongest H bond corresponds to the electrostatically enhanced N⁺–H...O[−] H bond (CP4), in agreement with the shortest distance.

Hirshfeld Surface Analysis. Hirshfeld surface (HS)⁴⁴ analysis and analysis of 2D fingerprint plots (2D FPs) derived from them⁴⁵ offer a unique means of exploration of intermolecular contacts within the crystal structures.^{46–50} One of the greatest benefits that comes from using these

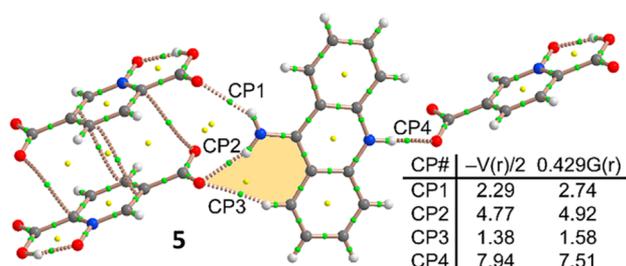


Figure 17. AIM distribution of bond and ring critical points (green and yellow spheres, respectively) and bond paths obtained for the H-bonded assembly of compound 5. The dissociation energies of the H bonds obtained using the $V(r)$ and $G(r)$ values at the bond CP are indicated (in kcal/mol) in the lower right corner.

techniques is the possibility for decomposition of the HS into regions dominated by specific contacts, which in turn allows one to calculate percentage contributions of various contact classes (e.g., $H\cdots H$ or $O\cdots H$) to the HS. In this way, supramolecular environments of molecules can be systematically compared to each other in quantitative terms. This approach proved to be particularly helpful in the identification of substituent, counterion, and (de)protonation effects on the HSs of related compounds.^{51–54}

From the viewpoint of HS analysis of supramolecular interactions, a very interesting feature of compounds 1–5 is that they combine two aspects of a proton transfer. As already indicated in previous sections, these compounds belong to the salt class (with the exception of 5, which should be referred to as an “ionic cocrystal”), since all feature intermolecular proton transfer. Additionally, the presence of a carboxylate group next to the *N*-oxide creates a possibility for an *intramolecular* transfer of protons between oxygen acceptors, giving rise to the tautomerism phenomenon. Accordingly, HS analysis should target and assess the effect both of these proton transfer phenomena on the properties of HSs. To this end, we have drawn d_{norm} -mapped HSs of Hpydco[−] (1–4, 5A, 6) and H₂pydco (5B) species (Figure S9, Supporting Information) as well as calculated contact-decomposed 2D FPs (Figures S10–S15, Supporting Information). In Table 2 are given percentage contributions of respective contacts for each HS.

Table 2. Percentage Contributions of Intermolecular Contacts to the HSs of 1–5

entry	1	2	3A	3B	4a	4b	5
O \cdots H	42.3	47.4	45.3	46.7	51.0	51.6	49.5
H \cdots H	16.4	15.1	21.9	21.0	14.5	15.3	22.8
C \cdots H	9.6	11.1	9.3	9.3	6.9	12.8	10.1
C \cdots O	12.3	8.1	11.1	10.8	4.4	10.6	4.5
C \cdots C	3.2	5.9	5.6	5.6	13.8	2.9	6.0
N \cdots H	5.7	4.5	2.1	2.1	3.0	0.7	3.2
O \cdots O	3.7	3.5	0.7	0.6	1.8	2.2	2.3
N \cdots C	5.2	3.0	2.2	2.2	4.4	2.0	0.3

A general description of features of HSs and 2D FPs of 1–5 is provided in the Supporting Information, but here we will explore in depth specific HS properties of 3A,B, 4, and 5.

We will start from the analysis of protonation-dependent supramolecular roles that are played by Hpydco[−] and H₂pydco species. 4 is an example of an ionic cocrystal, since it involves both a Hpydco[−] anionic (4a) and H₂pydco (4b) neutral molecule, apart from the phenanthroline cation. Note that

small letters are used here to distinguish Hpydco[−] and H₂pydco molecules. The presence of two molecules of different protonation states in one crystal lattice allows reliable side by side comparisons of their contact contributions to the HS (Table 2). Indeed, quite strikingly, if one compares the percentage of O \cdots H contacts, it turns out that the protonation state does not significantly affect the relative contribution of this contact (51.0% vs 51.6% for 4a,b, respectively). In fact, the largest differences in contact contributions due to protonation emerge not in contacts primarily associated with strong hydrogen bonding (O \cdots H and N \cdots H) but in those regarded rather as supporting contacts. For instance, a dramatic difference is noted for C \cdots C contacts (13.8% vs 2.9% for 4a,b, respectively), C \cdots H contacts (6.9% vs 12.8% for 4a,b, respectively), and C \cdots O contacts (4.4% vs 10.6% for 4a,b, respectively). The location of these contacts is presented in Figure 18a. These differences in contact contributions are easy

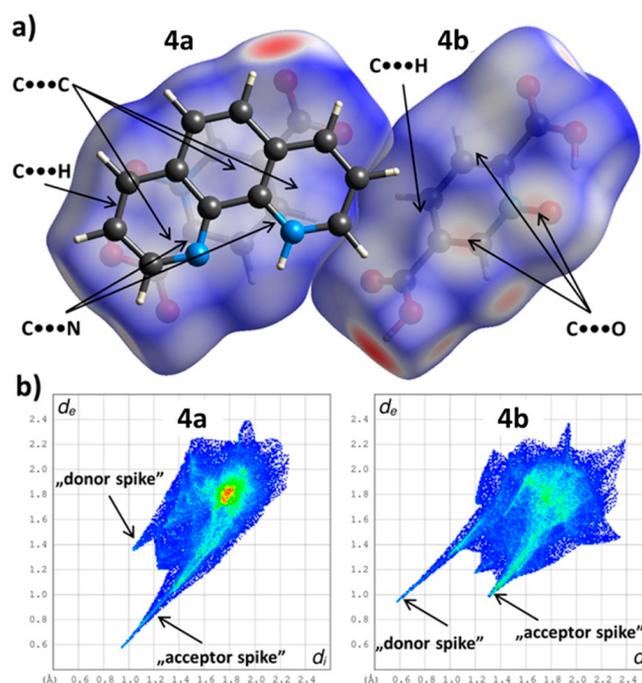


Figure 18. (a) Semitransparent HSs mapped with the d_{norm} function for 4a (left) and 4b (right). The most important intermolecular interactions are indicated with arrows; all red areas that are not described on the picture indicate O \cdots H contacts. (b) 2D FPs of 4a (left) and 4b (right).

to understand if one appreciates the various structural roles played by Hpydco[−] and H₂pydco molecules. As shown in Figure 10 in Description of the Crystal Structures, Hpydco[−] (4a) participates in extensive charge-assisted π – π stacking with phenanthroline cation, which is imposing a high contribution of C \cdots O and C \cdots C contacts. Note that the former contacts arise from the interaction of carboxylic groups with the π -conjugated plane of aromatic cations. On the other hand, π – π stacking with such a large aromatic molecule effectively shields Hpydco[−] from C–H \cdots π interactions, which is clearly reflected in a diminished share of C \cdots H contacts for 4a.

As seen in Figure 18b, 2D FPs of 4a,b are largely different in both occupied area and overall shape. Strong hydrogen bonding is reflected in so-called “spikes”, pointing to the

lower left side of each plot. The upper spike corresponds to the hydrogen bond donor sites that form C–H···O, N–H···O, and O–H···O interactions, while the lower spike represents all O-accepting sites of these interactions, as determined from contact-decomposed FP plots (Figures S14 and S15 in the Supporting Information). It is clear that despite essentially the same contribution of O···H contacts (51.0% vs 51.6% for 3A,B, respectively) the protonation-dependent differences in 2D FP properties are outstandingly sharp.

For a broader perspective, the above observations regarding O···H contacts should be put into the context of the other systems for which the protonation-dependent HS properties have been investigated. For example, we have very recently found that protonation of melamine is strongly correlated with a higher participation of strong hydrogen bonding contacts to oxygen acceptors (O···H). Indeed, monoprotonated melaminium species (Hme⁺) were found to feature a 50–100% increase in O···H contacts vs nonprotonated melamine and at least a 4-fold increase in O···H contacts for biprotonated melaminium species (H₂me²⁺), again in comparison to neutral melamine.⁵³ Clearly, melamine is a basic compound and hence a more “apples to apples” comparison of Hpydco[−] and H₂pydco contact contributions necessitates consideration of acidic species. To the best of our knowledge, the only study that targeted the effect of deprotonation on the HS and HS-related properties was that of Bialek et al.⁵⁴ In this study, no significant change in percentage contribution of O···H contacts to the HS has been identified for naphthalene-1,5-diphosphonic acid with different phosphonic species (−PO₃H₂ and −PO₃H[−]), which is accordance with our results.

When all these considerations are taken together, the observed differences in quantitative HS properties of Hpydco[−] and H₂pydco species are predominantly reflected in shares of secondary contacts such as C···C, C···O, and C···H, rather than in contact contributions of strong interactions (O···H, N···H). This is because acid molecules in the deprotonated state experience approximately the same amount of strong hydrogen bonding as in the neutral state but engage in a different set of weaker contacts.

Having established the effect of protonation state of pydco species on HSs and their properties, we moved to the investigation of the intramolecular proton transfer on the HSs of molecules. To the best of our knowledge, this issue has not been investigated thus far. For this reason, it is not even known whether HS analysis is able to reliably pinpoint differences between two tautomers. Indeed, proton migration is an intramolecular event, which suggests that its effect on the HS properties (which in general, characterizes intermolecular contacts) of tautomers should not be profound. On the other hand, intramolecular proton transfer may be assisted by an overall change in supramolecular environment of a molecule, consequently leading to e.g. a subtly different set of interactions experienced by the molecule.

In order to explore the capability of HS analysis to investigate intramolecular proton transfer, we also looked at the HS properties of polymorphic compounds 3A,B, since they differ in the proton localization: in 3A the proton is localized at the carboxylic oxygen atom, while in 3B it occupies the central position between the oxygen atoms of carboxylate and N-oxide.

An inspection of d_{norm} -mapped HSs (Figure 19a) drawn for Hpydco[−] units in 3A,B shows that they closely resemble each other; their main feature is the presence of two red spots that

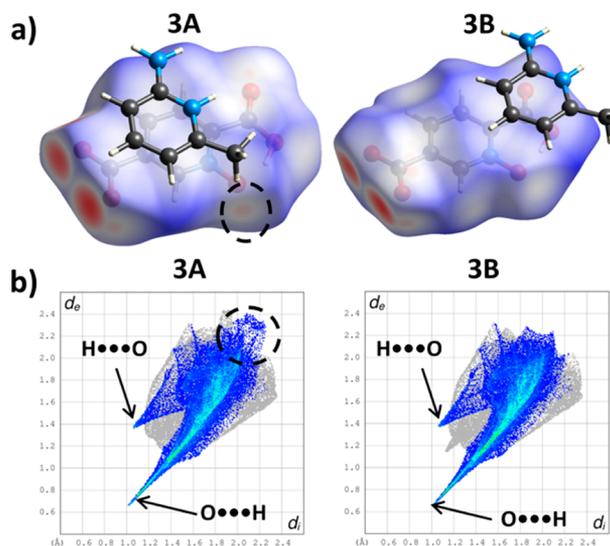


Figure 19. (a) Semitransparent HS mapped with the d_{norm} function for polymorphs 3A (left) and 3B (right). (b) Contact-decomposed (H···O) 2D FPs of 3A,B.

correspond to strong N–H···O contacts, contributing to HB R₂²(8) ring motifs. There is, however, a very subtle difference in the environment of the N-oxide functional group: in the case of 3A one sees a faint red area, whereas this is not present in 3B (Figure 19a). This contact originates from weak C_{ar}–H···O HBs between (H2a-6mpy)⁺ and Hpydco[−]. Contact-decomposed (H···O) 2D FPs of 3A,B show differences only in regions of high $d_1 + d_2$ values, i.e. where weak, distant contacts are reflected (denoted as a dashed oval in Figure 19b), highlighting the close similarity of supramolecular environments of Hpydco[−] species.

Contact contributions to the HS characterizing 3A,B are given in Table 2. It is apparent that values of percentage shares of contacts to the HS are quite similar, with the biggest difference of 0.9% visible in H···H contacts (21.9% vs 21.0% for 3A,B, respectively). This raises the question of whether slightly different contact contributions to the HS of 3A,B originate from their polymorphic structures or rather from the intramolecular shift of the hydrogen atom itself. In order to explore this issue, we have calculated contact contributions to the HS for specially modified crystal structures of 3A,B. The applied modification consists of the shifting of a hydrogen atom that resides between the carboxylic oxygen atom and N-oxide, so that a competing tautomer is obtained. More specifically, in the case of 3A the hydrogen atom has been placed at the central position between oxygen atoms (3A-mod), while for 3B the hydrogen atom has been placed near the carboxylic oxygen atom (3B-mod). As shown in Table S3 in the Supporting Information, the position of the mobile hydrogen atom does not have any significant effect on contacts that contribute to HS properties, on the basis of a comparison of 3A-mod and 3B-mod to 3A,B, respectively. Consequently, minor differences in contact landscapes of 3A,B (Table 2) can be ascribed to various crystal environments, but not to intramolecular proton transfer (tautomerization). In light of these results it is suggested that HS analysis is not a sensitive tool for the differentiation of tautomeric structures.

Concluding Remarks. In summary, we prepared and characterized a series of acid–base compounds involving Hpydco[−] as the anion and several N-containing aromatic bases

as cations. An important observation is that the hydrogen-bonded motif $R_2^2(8)$ is present in all crystals structures that involve the α -aminopyridinium moiety (compounds **1**, **2**, and **3A,B**). The use of large polycyclic organic bases (phenanthroline and 9-aminoacridine, compounds **4** and **5**) resulted in the formation of hydrogen-bond networks devoid of this motif, which however has been replaced by extensive π - π -stacking and C-H \cdots π interactions. Accordingly, the presence of different kinds of cations predetermines how Hpydco⁻ will participate in the crystal packing.

We have computed the interaction energies of the individual H-bonding interactions observed in the solid state of compounds **1**–**5**, which led us to several interesting conclusions. One is that positively charged melaminium species in **1** form antielectrostatic N–H \cdots N H bonds which are energetically favorable (around 4.0 kcal/mol). The other is that charge-assisted hydrogen bonding does not always offer an advantage over “noncharged” hydrogen bonds. For example, the N⁺–H \cdots O bond is stronger than the exocyclic N–H \cdots O bond in **2**, whereas the opposite relationship is observed in **1**, which suggests stronger delocalization in the latter. One more example of that is **4**, whose strongest interaction in the hydrogen-bonding net corresponds to the O–H \cdots O H bond rather than to the electrostatically “enhanced” N⁺–H \cdots O H bond.

Taken together, the energetic study might be useful to rationalize cocrystallization modes of pyridine-2,5-dicarboxylic acid *N*-oxide derivatives.

The effect of the protonation state of pydco species on HSs 2D FP properties has been investigated. It is observed that differences in HS properties of Hpydco⁻ and H₂pydco species, in quantitative terms, are mainly reflected in shares of secondary contacts such as C \cdots C, C \cdots O, and C \cdots H, rather than in contact contributions of strong interactions (O \cdots H, N \cdots H). On the other hand, the biggest effect on the appearance of 2D FPs of Hpydco⁻ and H₂pydco species is different sets (either primarily accepting or donating) of O \cdots H contacts to these molecules.

We have also investigated the usefulness of HS analysis for a comparison of polymorphic structures containing species with intramolecular proton transfer. We found the bulk of qualitative differences in HS and 2D FP properties are only due to various crystal environments and are not due to intramolecular proton transfer. Accordingly, we suggest that HS analysis is not a sensitive tool for the differentiation of tautomeric structures in similar crystal environments.

■ ASSOCIATED CONTENT

SI Supporting Information

The Supporting Information is available free of charge at <https://pubs.acs.org/doi/10.1021/acs.cgd.9b01475>.

Selected bond lengths (Å) and valence angles (deg) for **1**–**5**, selected hydrogen bond geometries, infrared spectra, additional packing figures, Cartesian coordinates, and Hirshfeld surface analysis (PDF)

Accession Codes

CCDC 1554218–1554220, 1554228–1554229, and 1554231 contain the supplementary crystallographic data for this paper. These data can be obtained free of charge via www.ccdc.cam.ac.uk/data_request/cif, or by emailing data_request@ccdc.cam.ac.uk, or by contacting The Cambridge Crystallographic

Data Centre, 12 Union Road, Cambridge CB2 1EZ, UK; fax: +44 1223 336033.

■ AUTHOR INFORMATION

Corresponding Authors

Masoud Mirzaei – Department of Chemistry, Faculty of Science, Ferdowsi University of Mashhad, Mashhad, Iran; orcid.org/0000-0002-7256-4601; Email: mirzaeesh@um.ac.ir; Fax: +98 38796416

Jan K. Zareba – Advanced Materials Engineering and Modelling Group, Wrocław University of Science and Technology, 50370 Wrocław, Poland; orcid.org/0000-0001-6117-6876; Email: jan.zareba@pwr.edu.pl

Antonio Frontera – Department of Chemistry, Universitat de les Illes Balears, 07122 Palma, Balears, Spain; orcid.org/0000-0001-7840-2139; Email: toni.frontera@uib.es; Fax: +34 971 173426

Authors

Fereshteh Sadeghi – Department of Chemistry, Faculty of Science, Ferdowsi University of Mashhad, Mashhad, Iran

Krešimir Molčanov – Rudjer Bošković Institute, HR-10000 Zagreb, Croatia

Rosa M. Gomila – Department of Chemistry, Universitat de les Illes Balears, 07122 Palma, Balears, Spain

Complete contact information is available at: <https://pubs.acs.org/10.1021/acs.cgd.9b01475>

Author Contributions

The manuscript was written through contributions of all authors. All authors have given approval to the final version of the manuscript.

Notes

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