

Crystal Structure and Hirshfeld Surface Analysis Of Diethyl (6-methyl-2-pyridyl)aminoethylenemalonate

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Abstract

The title compound, diethyl (6-methyl-2-pyridyl)aminoethylenemalonate (**1**), crystallises in the monoclinic space group $P2_1/c$ (No.14). The unit cell parameters are $a = 10.5657(7)$ Å, $b = 9.1784(5)$ Å, $c = 14.5681(7)$ Å, $\beta = 101.636(6)^\circ$, $Z' = 1$ and $Z = 4$ at 150 K. The extended structure forms approximately orthogonal columns of stacked molecules. All bond lengths and angles are unremarkable. No disorder, twinning or co-crystallised solvent is present in the structure. An intramolecular hydrogen bond exists between the enamine nitrogen and carbonyl oxygen. Hirshfeld surface analysis reveals a short contact between a carbonyl oxygen and neighbouring aryl hydrogen, as well as a carbonyl-carbonyl interaction.

Keywords

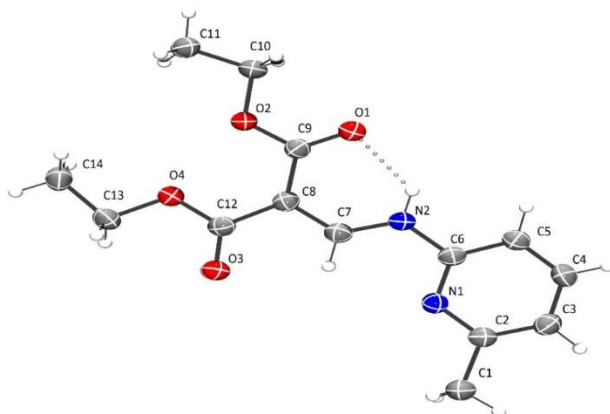
Organic structure, intramolecular hydrogen bonding, Hirshfeld surface analysis, fingerprint plots, X-ray crystallography.

Index Abstract

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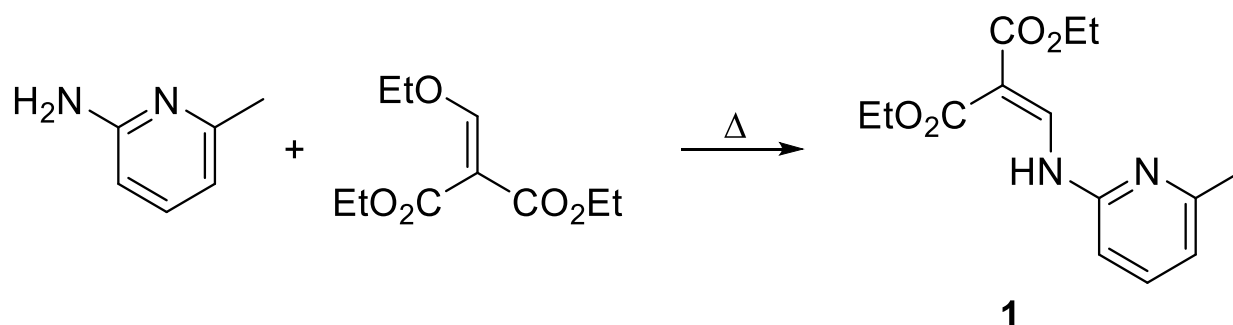
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The X-ray structure of diethyl (6-methyl-2-pyridyl)aminoethylenemalonate (**1**), an intermediate in the synthesis of ethyl nalidixate, is reported, and Hirshfeld surface analysis employed to identify intermolecular interactions within the structure.



Introduction

Initially reported in 1962 [1], the title compound diethyl (6-methyl-2-pyridyl)aminoethylenemalonate (**1**) is commonly prepared as an intermediate in the synthesis of ethyl nalidixate in undergraduate medicinal chemistry laboratories. Ethyl nalidixate has since become ubiquitous in medicinal chemistry as a lead compound for the development of quinolone antibiotics [2]. Synthesis of **1** entails a condensation reaction between 2-aminopicoline and diethyl ethoxymethylenemalonate (Scheme 1). Despite its role as a key characterisation technique for inorganic and organometallic compounds, single crystal X-ray diffraction (SC-XRD) is much less routinely employed for the characterisation of neutral organic compounds, and the solid-state structure of **1** has not been previously reported.



Scheme 1. Synthesis and molecular structure of **1**

Hirshfeld surface analysis [3] has recently become prevalent for describing interactions within crystals, and provides a method to view molecules as ‘organic wholes’ [4]. It offers a tool for visualising interactions in crystal structures, allowing for comparison between polymorphs and chemically similar structures [5]. By mapping the distance from the nearest external nucleus (d_e) onto the Hirshfeld surface, represented as colour contours on the generated 3D visualisations, we can identify points of intermolecular interaction. Plotting d_e against d_i (the distance to the nearest internal nucleus) gives 2D-fingerprint plots which allow more detailed analysis of these interactions.

Experimental

NMR spectra were recorded on a JEOL Eclipse 400 MHz NMR spectrometer (399.60 MHz for ^1H and 100.48 MHz for ^{13}C). LC/MS was carried out using a Waters Acquity UPLC (Waters Corp, Milford, MA, USA), coupled to a Waters Xevo G2-XS Q-TOF mass spectrometer (Waters Corp, Milford, MA, USA). Melting point measurements are uncorrected. Yield and spectra were in close agreement with literature values [2].

Synthesis of **1**

2-aminopicoline (2.00 g, 18.49 mmol) and diethyl ethoxymethylenemalonate (4.00 g, 18.49 mmol) were combined and heated at 110 °C for 90 minutes with stirring. The precipitate formed on cooling was recrystallized from ethanol to afford **1** as a white powder. Yield: 4.11 g, 80%.

^1H -NMR (400 MHz, CDCl_3): δ 11.04 (1H, d br, J = 12.58 Hz, NH), 9.25 (1H, d, J = 13.04 Hz, C=C-H), 7.51 (1H, t, J = 7.78 Hz, Ar-H), 6.86 (1H, t, J = 7.55, Hz, Ar-H), 6.64 (1H, t, Ar-H), 4.27(4H, dq, J = 19.06 Hz, 7.08 Hz, OC- H_2), 2.48 (3H, s, Ar-C- H_3), 1.34 (3H, m, C- CH_3).

^{13}C -NMR(100 MHz, CDCl_3): δ 14.28 (s), 14.40 (s), 24.29 (s), 60.14 (s), 60.48 (s), 94.99 (s), 108.61 (s), 119.08 (s), 138.68 (s), 149.89 (s), 150.08 (s), 165.69 (s)

LC/MS $[\text{M}+\text{H}]^+$: Calc. for $\text{C}_{14}\text{H}_{19}\text{N}_2\text{O}_4$ = 279; found 279.

Melting point (ethanol): 104-106° C.

SC-XRD quality crystals were obtained by slow evaporation of a solution of **1** in tetrahydrofuran over 3 days.

X-Ray Crystallography

Crystallographic measurements were performed at 150 K using an Oxford Xcalibur Gemini diffractometer with a Sapphire 3 CCD plate (graphite-monochromated Cu K α radiation, $\lambda = 1.54184$ Å). A crystal of suitable size and quality was selected, coated with Fomblin® Y oil, and mounted onto a nylon loop. The Numerical absorption correction was based on gaussian integration over a multifaceted crystal model[6]. Empirical absorption correction, using spherical harmonics, implemented in SCALE3. The ABSPACK scaling algorithm was applied for absorption correction[7]. Cell refinement, data collection and data reduction were performed using Rigaku CrysAlisPro 1.171.39.46[8]. The crystal density was not measured.

Refinement

Using Olex2 [9], the structure was solved with the ShelXT structure solution program using intrinsic phasing and refined with the ShelXL refinement package using Least Squares minimisation [10, 11]. All hydrogen atoms were geometrically placed and refined using the riding model approximation. Methyl groups were refined as a rotating group. Structural and refinement parameters are given in table 1. Bond lengths, angles and atomic displacement parameters are given in the supplementary information.

Table 1. Crystal data and structure refinement of 1

CCDC Deposition Number	1877868
Empirical formula	C ₁₄ H ₁₈ N ₂ O ₄
Formula weight	278.30
Temperature/K	150.01(10)
Crystal system	Monoclinic
Space group	P2 ₁ /c
a/Å	10.5657(7)
b/Å	9.1784(5)
c/Å	14.5681(7)
α /°	90
β /°	101.636(6)
γ /°	90
Volume/Å ³	1383.73(14)
Z	4
$\rho_{\text{calc}}/\text{cm}^3$	1.336
μ/mm^{-1}	0.819
F(000)	592.0
Crystal size/mm	0.413 × 0.214 × 0.122
Radiation	Cu K α ($\lambda = 1.54184$)
2 θ range for data collection/°	8.544 to 133.168
Index ranges	-11 ≤ h ≤ 12, -6 ≤ k ≤ 10, -12 ≤ l ≤ 17
Reflections collected	3693
Independent reflections	2407 [$R_{\text{int}} = 0.0293$, $R_{\text{sigma}} = 0.0523$]
Data/restraints/parameters	2407/0/184
Goodness-of-fit on F^2	1.098
Final R indexes [$ I \geq 2\sigma(I)$]	$R_1 = 0.0726$, $wR_2 = 0.1949$
Final R indexes [all data]	$R_1 = 0.0902$, $wR_2 = 0.2121$
Largest diff. peak/hole / e Å ⁻³	0.37/-0.26

Structural Analysis

Bond parameters were measured using Mercury 3.10 [12–14]. Hirshfeld surface analysis was performed using CrystalExplorer [15]. The Hirshfeld surface was mapped using an isovalue of 0.5. Red contours indicate a contact less than the sum of the Van Der Waals radii of the respective elements. Blue and white contours indicate that the nearest external atom is at a distance greater than or equal to the sum of the Van Der Waals radii respectively from the atomic co-ordinate. As hydrogen atoms have been placed geometrically, e.s.d. are not included in quoted bond lengths.

Results and Discussion

Asymmetric Unit

1 crystallises in the monoclinic space group $P2_1/c$ (No.14). The asymmetric unit consists of one molecule of **1** (Figure 1) and the unit cell consists of four molecules of **1**.

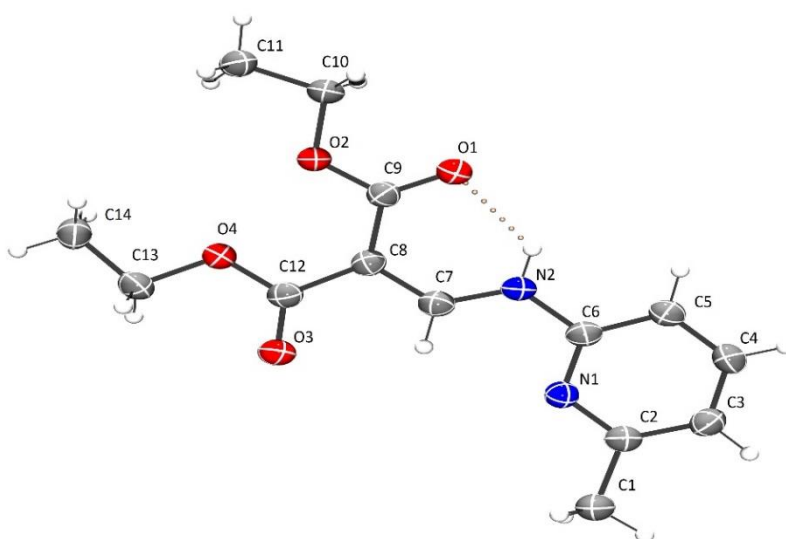


Figure 1. Representation of the solid-state structure of **1**. Thermal ellipsoids are shown at 50% probability. Hydrogen atom labels are omitted for clarity. The O1-N2 hydrogen bond is shown as a tan dashed line.

The molecular configuration of **1** is essentially planar in the solid state, with C4 and C14 having the largest deviation from the plane at 0.117 Å and 0.129 Å respectively. This planarity is supported by a hydrogen bonding interaction between O1 and N2, which forms a six-membered ring (O1-C9-C8-C7-N2-H2) and locks the conformation of the ester groups. The strongly downfield shift of the enamine proton in the ^1H NMR spectrum (11.04 ppm) suggests that this hydrogen bonding interaction is maintained in solution. No disorder, twinning or co-crystallised solvent is present in the structure. All bond lengths and angles are unremarkable. The bond lengths and angles in the heterocyclic component of **1** are similar to pyridine [16].

Intermolecular Interactions

Molecules are stacked in a head-to-tail fashion (Figure 2). The interplanar distance between molecules is 3.396 Å. Stacked molecules are fully co-planar, with an angle of 0.00° between them. These planes are offset by 2.84 Å. The resulting columns of stacked molecules are approximately orthogonal to one another with an angle of 86° between them.

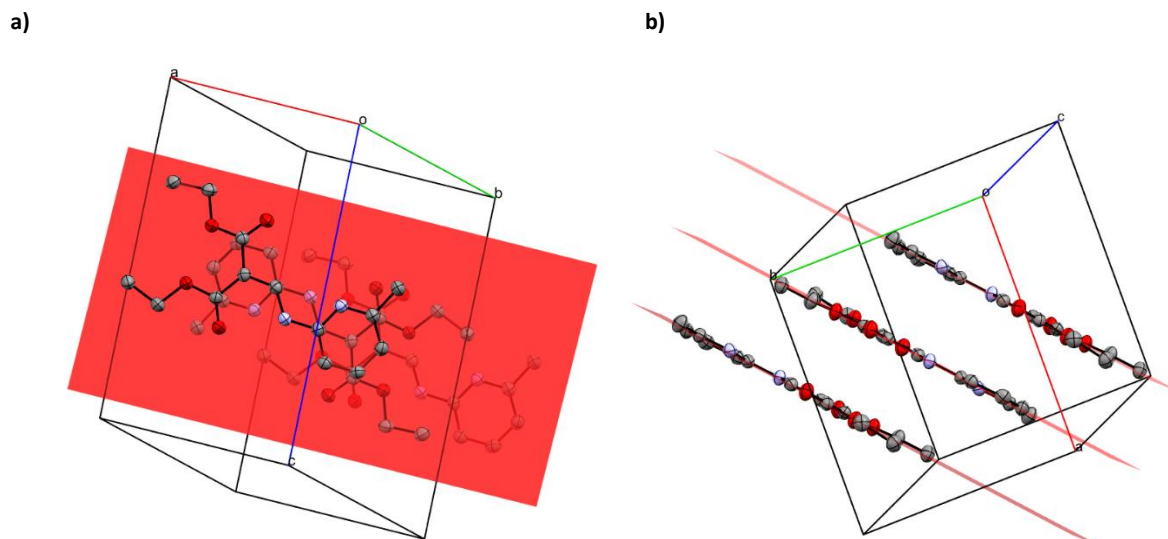


Figure 2. a) Parallel view of the asymmetric unit of **1** with stacked molecule above and below it. b) Perpendicular view of the asymmetric unit of **1** with stacked molecule above and below it. The molecule planes and cell axes are shown. Atom labels and hydrogen atoms omitted for clarity.

Database Survey

Eleven crystal structures possessing a 2-(2,2-Dicarbalkoxyvinylamino)-pyridine substructure exist in the Cambridge Structural Database (CSD, Version 5.39, November 2017). The most comparable to **1** is 2-(N-(2,2-Diformylethenyl)amino)pyridine (CCDC:297480, Figure 3). Similarly to **1**, a hydrogen bond interaction exists between the enamine N and a carbonyl O. The supramolecular structure is similar to **1**, in that the molecule is planar, with a shorter intermolecular stacking distance (3.389 Å).

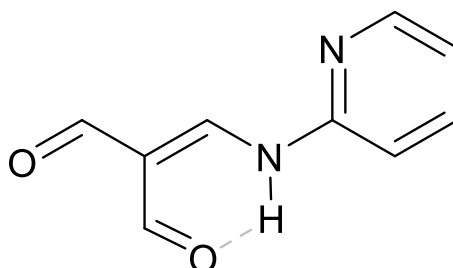


Figure 3. Structure of 2-(N-(2,2-Diformylethenyl)amino)pyridine (CCDC:297480), showing the intramolecular hydrogen bond in grey.

Hirshfeld Surface Analysis

The Hirshfeld analysis reveals a short contact between O3 and H5 of a neighbouring molecule of 2.402 Å, demonstrated by red contours on the Hirshfeld surface at these atoms (Figure 4). A fairly short contact is also identified between C9 and C12 (3.444 Å). This is likely the result of a carbonyl-carbonyl stacking interaction. Such interactions have been found in other small molecules and the secondary structures of proteins, and are the result of $n \rightarrow \pi^*$ and $\pi \rightarrow \pi^*$ interactions [17]. There is also a short contract between H3 and C11 (2.817 Å).

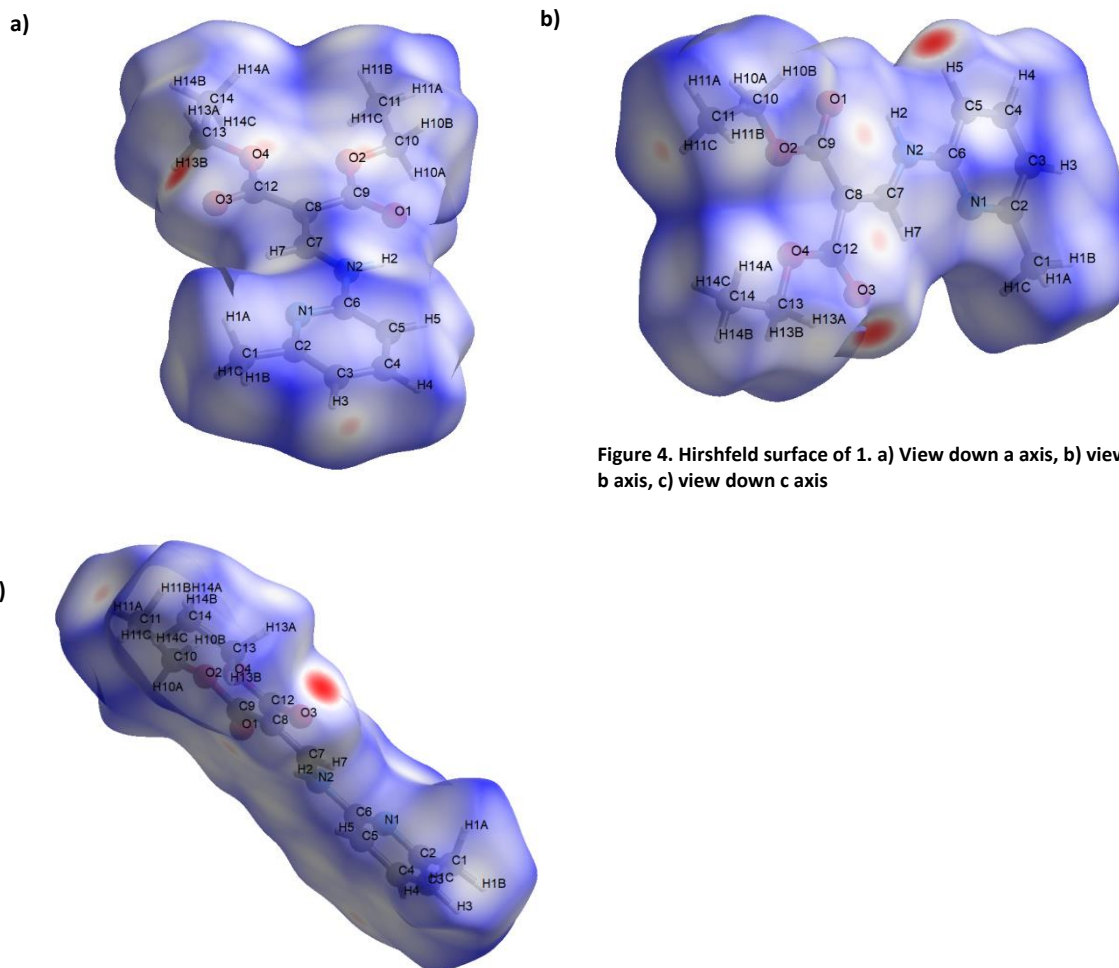


Figure 4. Hirshfeld surface of 1. a) View down a axis, b) view down b axis, c) view down c axis

The fingerprint plots (Figure 5) show that the predominant intermolecular interaction is H-H interactions; these account for 60% of the Hirshfeld surface (Figure 5 c). These interactions are presumed to be primarily the result of attractive dispersion forces [18]. C-C interactions, primarily the result of stacking interactions, also contribute to the Hirshfeld surface (Figure 5 b).

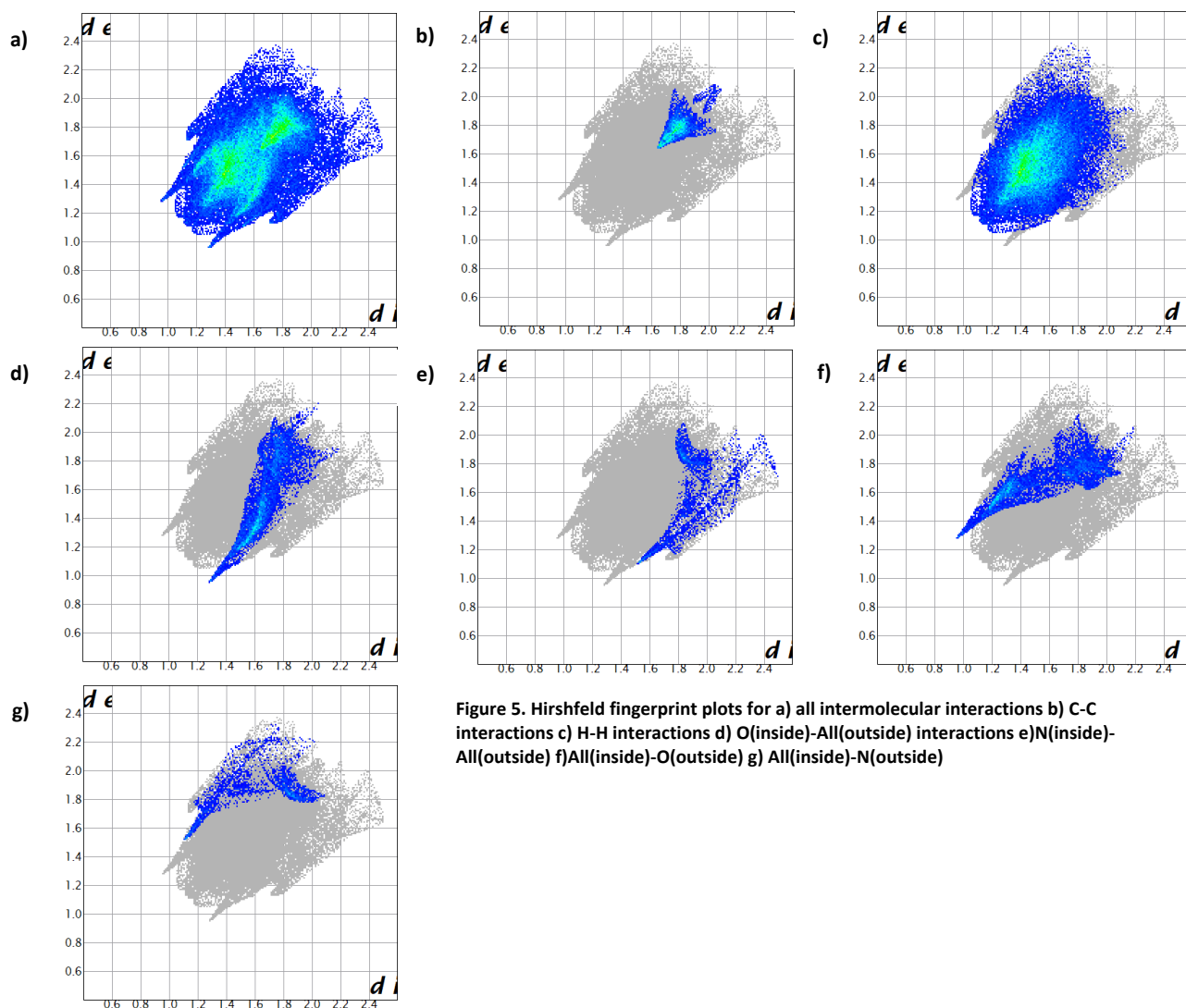


Figure 5. Hirshfeld fingerprint plots for a) all intermolecular interactions b) C-C interactions c) H-H interactions d) O(inside)-All(outside) interactions e) N(inside)-All(outside) f) All(inside)-O(outside) g) All(inside)-N(outside)

Conclusions

Here, we have presented the solid-state structural analysis of a common organic small molecule. Intramolecular H-bonding is present in the structure, and Hirshfeld surface analysis has been used to reveal intermolecular interactions that would otherwise have been difficult to elucidate.

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