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

Omar Coughlin<sup>1</sup>, Nathan De Bruyn<sup>2</sup>, David P. A. Kilgour<sup>1</sup>, Sophie L. Benjamin<sup>\*1</sup>

- 1. Department of Chemistry and Forensics, Nottingham Trent University, Clifton Lane, Nottingham NG11 8NS, United Kingdom
- 2. School of Chemistry, The University of Manchester, Oxford Road, Manchester M13 9PL, United Kingdom
  - \* email: sophie.benjamin@ntu.ac.uk

# Abstract

The title compound, diethyl (6-methyl-2-pyridyl)aminoethylenemalonate (1), crystallises in in the monoclinic space group P2<sub>1</sub>/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)°, 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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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 <sup>1</sup>H and 100.48 MHz for <sup>13</sup>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%.

<sup>1</sup>H-NMR (400 MHz, CDCl<sub>3</sub>):  $\delta$  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*<sub>2</sub>), 2.48 (3H, s, Ar-C-*H*<sub>3</sub>), 1.34 (3H, m, C-C*H*<sub>3</sub>).

<sup>13</sup>C-NMR(100 MHz, CDCl<sub>3</sub>): 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)

 $LC/MS [M+H]^+$ : Calc. for  $C_{14}H_{19}N_2O_4 = 279$ ; 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 Xcaliubur Gemini diffractometer with a Sapphire 3 CCD plate (graphite-monochromated Cu K $\alpha$  radiation,  $\lambda$  = 1.54184 Å). A crystal of suitable size and quality was selected, coated with Fomblin<sup>®</sup> 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.

CCDC Deposition Number	1877868
Empirical formula	C <sub>14</sub> H <sub>18</sub> N <sub>2</sub> O <sub>4</sub>
Formula weight	278.30
Temperature/K	150.01(10)
Crystal system	Monoclinic
Space group	P21/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/ų	1383.73(14)
Z	4
ρ <sub>calc</sub> g/cm <sup>3</sup>	1.336
µ/mm⁻¹	0.819
F(000)	592.0
Crystal size/mm	0.413 × 0.214 × 0.122
Radiation	Cu Kα (λ = 1.54184)
20 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 <sub>int</sub> = 0.0293, R <sub>sigma</sub> = 0.0523]
Data/restraints/parameters	2407/0/184
Goodness-of-fit on F <sup>2</sup>	1.098
Final R indexes [I>=2σ (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 <sup>-3</sup>	0.37/-0.26

#### Table 1. Crystal data and structure refinement of 1

### **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 <sup>1</sup>H 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).



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).

C5 C4

C1 H1B H1C H1A

C



# 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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