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Key indicators

Single-crystal X-ray study T = 293 KMean σ (C–C) = 0.004 Å R factor = 0.041 wR factor = 0.116 Data-to-parameter ratio = 12.2

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

Methyl 5-nitro-2-(phenylamino)pyridine-4-carboxylate

In the crystal structure of the title compound, $C_{13}H_{11}N_3O_4$, there are two independent molecules in the asymmetric unit. In each case, the methylcarboxylate group is twisted by about 90° relative to the rest of the molecule. The molecules are packed in the crystal structure through N $-H \cdots O$ hydrogen bonds.

Comment

Many pharmaceutical products consist of oxygen and/or nitrogen heterocycles, and the title compound, (I), is the result of a study to produce bis-heterocyclic compounds from pyridyl isocyanates. It crystallizes in the space group $P\overline{1}$ with two molecules in the asymmetric unit (Fig. 1). The phenyl and pyridine rings make a dihedral angle of 11.9 (2)° in one molecule and 11.9 (3)° in the other. In both molecules, the nitro group is almost coplanar with the pyridine ring, while the methylcarboxylate group is approximately perpendicular to this ring, due to steric hindrance; this is demonstrated by the selected torsion angles in Table 1. Bond lengths are all within the normal ranges (Allen *et al.*, 1987) and equivalent bond lengths in the two independent molecules are equal within their standard deviations. The molecules are packed in the crystal structure through N-H···O hydrogen bonds (Table 2).



Experimental

Methyl 5-nitro-2-(phenylamino)-4-pyridinecarboxylate, (I), was prepared by oxidative nucleophilic substitution (Finch *et al.*, 1980) of methyl 3-nitro-4-pyridinecarboxylate (Holt *et al.*, 2005) by sodio-formanilide and subsequent hydrolysis (Tjosaas & Fiksdahl, 2006). Crystals were grown by vapour diffusion of pentane into a solution in acetone in a refrigerator.

V = 1271.8 (5) Å³

 $D_x = 1.427 \text{ Mg m}^{-3}$ Mo $K\alpha$ radiation $\mu = 0.11 \text{ mm}^{-1}$ T = 293 (2) KBlock, light yellow

 $0.50 \times 0.40 \times 0.30 \text{ mm}$

Z = 4

Crystal data

C ₁₃ H ₁₁ N ₃ O ₄	
$M_r = 273.25$	
Triclinic, P1	
a = 9.990 (2) Å	
<i>b</i> = 10.913 (3) Å	
c = 12.6758 (19) Å	
$\alpha = 81.055 \ (17)^{\circ}$	
$\beta = 78.137 \ (16)^{\circ}$	
$\gamma = 70.86 \ (2)^{\circ}$	

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023

021

N13

N11

1014

Data collection

Enraf-Nonius CAD-4 diffractometer ω -2 θ scans Absorption correction: ψ scan (*ABSCALC* in *OSCAIL*; McArdle & Daly, 1999; North *et al.*, 1968) $T_{min} = 0.948, T_{max} = 0.968$

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.041$ $wR(F^2) = 0.116$ S = 0.954452 reflections 364 parameters H-atom parameters constrained 4900 measured reflections 4452 independent reflections 2078 reflections with $I > 2\sigma(I)$ $R_{int} = 0.018$ $\theta_{max} = 25.0^{\circ}$ 3 standard reflections frequency: 120 min

intensity decay: 1%

$$\begin{split} w &= 1/[\sigma^2(F_o^2) + (0.0576P)^2] \\ \text{where } P &= (F_o^2 + 2F_c^2)/3 \\ (\Delta/\sigma)_{\text{max}} &= 0.021 \\ \Delta\rho_{\text{max}} &= 0.19 \text{ e } \text{\AA}^{-3} \\ \Delta\rho_{\text{min}} &= -0.17 \text{ e } \text{\AA}^{-3} \\ \text{Extinction correction: } SHELXL97 \\ \text{Extinction coefficient: } 0.0074 (14) \end{split}$$

 Table 1

 Selected torsion angles (°).

O14-N13-C19-C110	178.7 (2)	C111-C110-C112-O12	90.4 (3)
C211-C210-C212-O22	-94.7 (3)	O24-N23-C29-C210	-172.3 (3)

Table 2

Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots \mathbf{A}$
$\begin{array}{c} \hline N21 - H21 \cdots O11 \\ N21 - H21 \cdots O11^{i} \\ N11 - H11 \cdots O21^{ii} \end{array}$	0.86	2.25	3.060 (3)	158
	0.86	2.62	3.107 (3)	117
	0.86	2.23	3.094 (3)	179

Symmetry codes: (i) -x, -y + 1, -z + 1; (ii) x, y - 1, z.

H atoms were positioned geometrically and refined using a riding model, with aromatic C-H = 0.93 Å, methyl C-H = 0.96 Å, N-H = 0.86 Å and $U_{\rm iso}({\rm H}) = 1.3U_{\rm eq}({\rm C},{\rm N})$ or $1.4U_{\rm eq}({\rm methyl}~{\rm C})$.

Data collection: *CAD-4-PC* Software (Enraf–Nonius, 1992); cell refinement: *CELDIM* in *CAD-4-PC* Software; data reduction: *XCAD4* (McArdle & Higgins, 1995); program(s) used to solve structure: *OSCAIL* (McArdle, 2005) and *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *OSCAIL* and *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEX* (McArdle, 1993) and *ORTEPIII* (Burnett & Johnson, 1996); software used to prepare material for publication: *OSCAIL*.



Figure 1

The asymmetric unit of (I), showing the atomic numbering scheme. Displacement ellipsoids are drawn at the 30% probability level.

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