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## Key indicators

Single-crystal X-ray study
$T=220 \mathrm{~K}$
Mean $\sigma(\mathrm{C}-\mathrm{C})=0.004 \AA$
$R$ factor $=0.039$
$w R$ factor $=0.109$
Data-to-parameter ratio $=13.1$

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.
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## 2-(2,4-Dichlorophenoxy)- N -(4,6-dimethyl-pyridin-2-yl)acetamide

The structure of the title compound, $\mathrm{C}_{15} \mathrm{H}_{14} \mathrm{Cl}_{2} \mathrm{~N}_{2} \mathrm{O}_{2}$, comprises an essentially flat molecule [dihedral angle of $3.64(9)^{\circ}$ between the two aromatic rings] with an intramolecular hydrogen-bonding interaction from the amine $\mathrm{N}-\mathrm{H}$ group to the phenoxy O atom. Molecules are arranged in lamellar sheets parallel to the (110) plane.

## Comment

A search of the April 2003 release of the Cambridge Structural Database (Allen, 2002) reveals that there are eight structures based on $N$-(4,6-dimethylpyridin-2-yl)carboxamide, all published by one research group. These structures are N -(4,6-dimethylpyridin-2-yl)-2-(3-nitrophenyl)acetamide (Rodier et al., 1986), $N$-(4,6-dimethylpyridin-2-yl)-2-(4-nitrophenyl)propionamide (Rodier, Robert \& Le Baut, 1990), (E)-$N$-(4,6-dimethylpyridin-2-yl)-3-phenylpropenamide hemihydrate (Rodier, Robert-Piessard \& Le Baut, 1990), N-(4,6-dimethylpyridin-2-yl)-2-furamide (Rodier et al., 1991), N-(4,6-dimethylpyridin-2-yl)-2-thiophenecarboxamide (Rodier, Cense et al., 1992), $N$-(4,6-dimethylpyridin-2-yl)(1-methylindol-2-yl)carboxamide (Rodier, Robert \& Le Baut, 1992), N-(4,6-di-methylpyridin-2-yl)-2-(3-nitrophenyl)acetamide (Rodier et al., 1993 ) and N -(4,6-dimethylpyridin-2-yl)-5-methylpyrazine-2carboxamide (Rodier et al., 1994). In a series of studies on the syntheses of additional derivatives of $N$-(4,6-dimethylpyridin-2yl)carboxamide and also $N$-(4,6-dimethylpyrimidin-2-yl)carboxamide as potential anti-inflammatory agents, we prepared the title compound, (I), and its structure is reported here.

(I)

The structure of (I) comprises an essentially flat molecule (Fig. 1) with an intramolecular hydrogen-bonding interaction from the amine $\mathrm{N}-\mathrm{H}$ group to the phenoxy O atom (Table 1). Molecules of (I) are arranged in lamellar sheets parallel to the (110) plane. The dihedral angle between the substituted phenyl and pyridyl rings is $3.64(9)^{\circ}$.

## Experimental

Four molar equivalents of oxalyl chloride ( $2.29 \mathrm{~g}, 18.1 \mathrm{mmol}$ ) were added dropwise to a stirred solution of (2,4-dichlorophenoxy) acetic acid $(1.0 \mathrm{~g}, 4.5 \mathrm{mmol})$ and a catalytic amount of dry pyridine (dried over KOH ) in 20 ml dry dichloromethane (dried over molecular sieves). After stirring for 30 min , the solvent was removed under

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Figure 1
The molecular configuration and atom-numbering scheme for the title compound, showing $50 \%$ probability displacement ellipsoids.
reduced pressure, yielding a yellow solution of (2,4-dichlorophenoxy)acetic chloride. This solution was then added dropwise to a stirred solution of 2-amino-4,6-dimethylpyridine ( $0.55 \mathrm{~g}, 4.5 \mathrm{mmol}$ ) and triethylamine ( $0.46 \mathrm{~g}, 4.5 \mathrm{mmol}$ ) in 20 ml dry dichloromethane. After 30 min , the solution was filtered and the solvent removed under reduced pressure, yielding a white powder. The product was purified using column chromatography $\left(\mathrm{SiO}_{2}\right)$ and collected in the eluted ethyl acetate fraction after initial chloroform elutions. The solid product was further washed with 30 ml cold ethanol to afford 0.82 g of (I) ( $81 \%$ ); m.p. $435-436 \mathrm{~K} ;{ }^{1} \mathrm{H}$ NMR $\left(\mathrm{CDCl}_{3}, 250 \mathrm{MHz}\right.$ ): $\delta$ (p.p.m.) $2.33\left(s, 3 H, \mathrm{CH}_{3}\right), 2.42\left(s, 3 \mathrm{H}, \mathrm{CH}_{3}\right), 4.62\left(s, 2 \mathrm{H}, \mathrm{CH}_{2}\right), 6.77(s, 1 \mathrm{H}, \mathrm{Ar}-$ H), $6.89(d, J=8.8 \mathrm{~Hz}, 1 \mathrm{H}, \mathrm{Ar}-\mathrm{H}), 7.33(d, J=8.8 \mathrm{~Hz}, 1 \mathrm{H}, \mathrm{Ar}-\mathrm{H}), 7.41$ $(d, J=2.5 \mathrm{~Hz}, 1 \mathrm{H}$, py-H), $7.87(s, 1 \mathrm{H}, \mathrm{py}-\mathrm{H}), 8.89(b s, 1 \mathrm{H}, \mathrm{NH}) ; m / z$ $325\left(\mathrm{MH}^{+}, 100 \%\right), 327\left(\mathrm{MH}^{+}, 75 \%\right)$. Crystals suitable for X-ray diffraction studies were grown from ethyl acetate.

## Crystal data

| $\mathrm{C}_{15} \mathrm{H}_{14} \mathrm{Cl}_{2} \mathrm{~N}_{2} \mathrm{O}_{2}$ | $Z=2$ |
| :--- | :--- |
| $M_{r}=325.18$ | $D_{x}=1.413 \mathrm{Mg} \mathrm{m}^{-3}$ |
| Triclinic, $P \overline{1}$ | Mo $K \alpha$ radiation |
| $a=7.445(5) \AA$ | Cell parameters from 22 |
| $b=8.790(6) \AA$ | reflections |
| $c=12.549(7) \AA$ | $\theta=15-16^{\circ}$ |
| $\alpha=89.28(4)^{\circ}$ | $\mu=0.43 \mathrm{~mm}^{-1}$ |
| $\beta=87.08(5)^{\circ}$ | $T=220(2) \mathrm{K}$ |
| $\gamma=68.74(4)^{\circ}$ | Prism, colourless |
| $V=764.4(9) \AA^{\circ}$ | $0.70 \times 0.47 \times 0.38 \mathrm{~mm}$ |
|  |  |
| Data collection |  |
| Stoe Stadi-4 diffractometer | $h=-8 \rightarrow 8$ |
| $\omega / \theta$ scans | $k=-10 \rightarrow 10$ |
| 3678 measured reflections | $l=0 \rightarrow 14$ |
| 2570 independent reflections | 3 standard reflections |
| 2032 reflections with $I>2 \sigma(I)$ | frequency: 60 min |
| $R_{\text {int }}=0.019$ | intensity decay: $3 \%$ |
| $\theta_{\text {max }}=25.0^{\circ}$ |  |

## Refinement

Refinement on $F^{2}$

$$
\begin{aligned}
& w=1 /[ \sigma^{2}\left(F_{o}^{2}\right)+(0.062 P)^{2} \\
&+0.1634 P] \\
& \text { where } P=\left(F_{o}{ }^{2}+2 F_{c}^{2}\right) / 3 \\
&(\Delta / \sigma)_{\max }<0.001 \\
& \Delta \rho_{\max }=0.20 \mathrm{e} \AA^{-3} \\
& \Delta \rho_{\min }=-0.30 \mathrm{e}^{-3}
\end{aligned}
$$

$w R\left(F^{2}\right)=0.109$
$S=1.05$
2570 reflections
196 parameters
H -atom parameters constrained

Table 1
Hydrogen-bonding geometry $\left(\AA{ }^{\circ}{ }^{\circ}\right)$.

| $D-\mathrm{H} \cdots A$ | $D-\mathrm{H}$ | $\mathrm{H} \cdots A$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :--- | :--- | :--- | :--- | :--- |
| $\mathrm{~N} 11-\mathrm{H} 11 \cdots \mathrm{O} 7$ | 0.87 | 2.07 | $2.552(3)$ | 114 |

All H atoms were included in the refinement at calculated positions, as riding atoms, with $\mathrm{N}-\mathrm{H}$ set to 0.87 A and $\mathrm{C}-\mathrm{H}$ set to 0.97 $\left(\mathrm{CH}_{3}\right), 0.98\left(\mathrm{CH}_{2}\right)$ or $0.94 \AA(\mathrm{Ar}-\mathrm{H})$; the isotropic displacement parameters were set at 1.25 times $U_{\text {eq }}$ of the carrier atom.

Data collection: DIF4 (Stoe \& Cie, 1990); cell refinement: DIF4; data reduction: REDU4 (Stoe \& Cie, 1990); program(s) used to solve structure: SHELXS 97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: PLATON97 (Spek, 1997); software used to prepare material for publication: SHELXL97.

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