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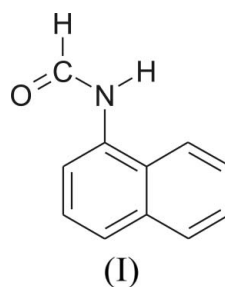
Key indicators

Single-crystal X-ray study
 $T = 173$ K
Mean $\sigma(\text{C}-\text{C}) = 0.002$ Å
 R factor = 0.030
 wR factor = 0.080
Data-to-parameter ratio = 10.1For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.*N*-(1-Naphthyl)formamide

In the title molecule, $\text{C}_{11}\text{H}_9\text{NO}$, all bond lengths and angles are normal. The crystal structure is stabilized by intermolecular $\text{N}-\text{H}\cdots\text{O}$ and $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds and $\pi-\pi$ stacking interactions.

Comment

N-Phenylformamides display a variety of packing patterns (Taylor & Kennard, 1982, 1983; Taylor *et al.*, 1983; 1984*a,b*; Leiserowitz & Tuval, 1978). In continuation of our systematic study of disubstituted *N*-phenylformamides in the crystalline state (Omondi *et al.*, 2005; Omondi, 2006), we present here the crystal structure of the title compound, (I).



In compound (I) (Fig. 1), all bond lengths and angles are normal (Allen *et al.*, 1987). The $\text{C}2-\text{C}1-\text{N}1-\text{C}11$ torsion angle is $53.0(2)^\circ$. The orientation of the formamide unit in (I) is similar to that observed in 2,6-disubstituted phenylformamides (Omondi *et al.*, 2005), but different from that found in *N*-2-naphthylacetamide (Haisa *et al.*, 1980). In the latter structure, the naphthyl and acetamide fragments are almost coplanar and form sheets of hydrogen-bonded molecules.

In the crystal structure of (I) (Fig. 2), $\text{N}-\text{H}\cdots\text{O}$ hydrogen bonds (Table 1) link the molecules into chains running along the short *a* axis. The molecules in the chains are related by translation; the naphthyl ring systems are stacked parallel to each other, with a relatively short $\text{C}g1\cdots\text{C}g2^{\text{iii}}$ distance of $3.759(1)$ Å, corresponding to $\pi-\pi$ intermolecular interactions [$\text{C}g1$ and $\text{C}g2$ are the centroids of the $\text{C}1-\text{C}5/\text{C}10$ and $\text{C}5-\text{C}10$ rings, respectively; symmetry code: (iii) $-1 + x, y, z$]. Weak intermolecular $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds (Table 1) contribute to the stability of the crystal packing.

Using the *OPIX* suite of programs (Gavezzotti, 2003), the lattice energy of (I) was calculated to be -118.36 kJ mol $^{-1}$. These calculations allowed us to estimate the quantitative contributions to this energy of the $\text{N}-\text{H}\cdots\text{O}$, $\pi-\pi$ and $\text{C}-\text{H}\cdots\text{O}$ intermolecular interactions as -49.97 , -14.40 and -13.96 kJ mol $^{-1}$, respectively.

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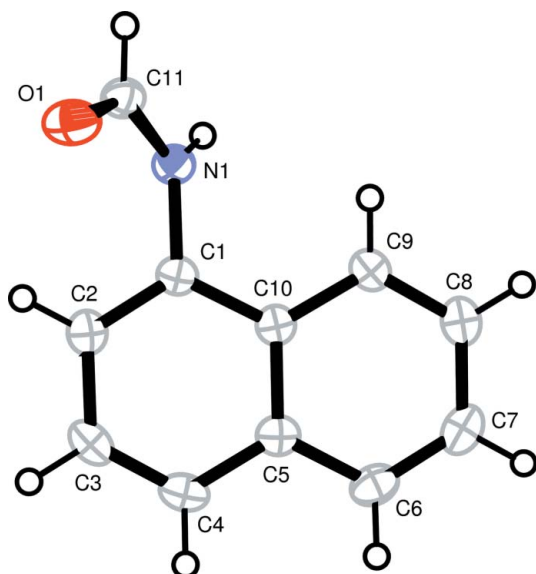


Figure 1
The molecular structure of (I), showing the atomic numbering scheme and 50% probability displacement ellipsoids.

Experimental

N-Naphthalenylformamide was synthesized following a known procedure (Ugi *et al.*, 1965). Commercially available 1-naphthylamine (Aldrich, purity > 95%) was heated in a tenfold excess of formic acid for a period of 15 h at 363 K. The excess formic acid was then removed under vacuum to give a purple solid, which was treated with dilute hydrochloric acid (0.1 M HCl, 10 ml) and ethyl acetate (60 ml). The organic layer was separated from the aqueous layer, dried over magnesium sulfate and filtered. Purple needle-shaped crystals of *N*-naphthyl-1-formamide were grown from the filtrate.

Crystal data

$C_{11}H_9NO$	$Z = 4$
$M_r = 171.19$	$D_x = 1.357 \text{ Mg m}^{-3}$
Orthorhombic, $P2_12_1$	Mo $K\alpha$ radiation
$a = 4.4639 (2) \text{ \AA}$	$\mu = 0.09 \text{ mm}^{-1}$
$b = 11.8463 (3) \text{ \AA}$	$T = 173 (2) \text{ K}$
$c = 15.8514 (4) \text{ \AA}$	Needle, purple
$V = 838.23 (5) \text{ \AA}^3$	$0.6 \times 0.09 \times 0.09 \text{ mm}$

Data collection

Bruker APEXII CCD area-detector diffractometer	14873 measured reflections
ω scans	1236 independent reflections
Absorption correction: multi-scan (SADABS; Sheldrick, 1996)	1097 reflections with $I > 2\sigma(I)$
$T_{\min} = 0.930$, $T_{\max} = 0.992$	$R_{\text{int}} = 0.034$
	$\theta_{\text{max}} = 28.4^\circ$

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0443P)^2 + 0.1116P]$
$R[F^2 > 2\sigma(F^2)] = 0.030$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.080$	$(\Delta/\sigma)_{\text{max}} = 0.006$
$S = 1.06$	$\Delta\rho_{\text{max}} = 0.16 \text{ e \AA}^{-3}$
1236 reflections	$\Delta\rho_{\text{min}} = -0.16 \text{ e \AA}^{-3}$
122 parameters	
H atoms treated by a mixture of independent and constrained refinement	

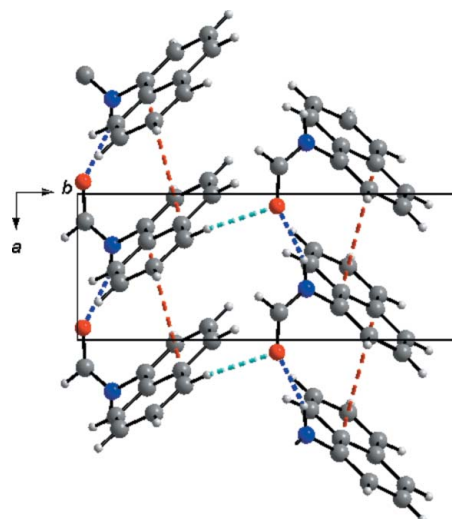


Figure 2
A portion of the crystal packing of (I), viewed down the c axis. Dashed lines denote hydrogen bonds and π - π stacking interactions.

Table 1

Hydrogen-bond geometry (\AA , $^\circ$).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$N1-H1\cdots O1^i$	0.85 (3)	2.00 (2)	2.7971 (19)	155.3 (18)
$C6-H6\cdots O1^{ii}$	0.95	2.50	3.436 (2)	170

Symmetry codes: (i) $x + 1, y, z$; (ii) $-x, y - \frac{1}{2}, -z + \frac{3}{2}$.

C-bound H atoms were positioned geometrically, with $C-H = 0.95 \text{ \AA}$, and allowed to ride on their parent atoms, with $U_{\text{iso}}(H) = 1.2U_{\text{eq}}(C)$. The N-bound atom H1 was located in a difference map and refined isotropically. In the absence of any significant anomalous scatterers in the molecule, the 826 Friedel pairs were merged before the final refinement.

Lattice energy calculations were performed using the *ZIPEPEC* module of the *OPIX* program suite (Gavezzotti, 2003).

Data collection: *SMART* (Bruker, 1997); cell refinement: *SAINT* (Bruker, 1997); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *PLATON* (Spek, 2003), *ORTEP-3* (Farrugia, 1997) and *DIAMOND* (Brandenburg, 2006); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

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