Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

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Key indicators

Single-crystal X-ray study T = 292 K Mean σ (C–C) = 0.005 Å R factor = 0.045 wR factor = 0.133 Data-to-parameter ratio = 8.1

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

In the title molecule, $C_{21}H_{15}NO$, the two ethynyl-bridged sixmembered rings are almost coplanar, making a dihedral angle of 3.3 (2)°. The dihedral angle between the two terminal phenyl rings is 62.9 (2)°. The crystal packing is stabilized by weak intermolecular $N-H\cdots O$ hydrogen bonds.

N-[4-(Phenylethynyl)phenyl]benzamide

Received 5 July 2005 Accepted 11 July 2005 Online 16 July 2005

Comment

The low-temperature luminescence of the title compound, (I), in non-polar solvents has been studied by Frederick & Liu (1999). In this paper, we report the crystal structure of (I). In the molecule (Fig. 1), the bond lengths and angles are in normal ranges (Allen *et al.*, 1987). The two ethynyl-bridged six-membered rings are almost coplanar, making a dihedral angle of 3.3 (2)°. The dihedral angle between the two terminal phenyl rings is $62.9 (2)^{\circ}$. The C12–N1–C15–C16 torsion angle is $178.8 (2)^{\circ}$. The crystal packing (Fig. 2) is stabilized by weak intermolecular N–H···O hydrogen bonds (Table 1), which link the molecules into chains along the *b* axis. The intermolecular N···O distance of 3.252 (3) Å is slightly longer than that in benzanilide [3.141 (4) Å; Bowes *et al.*, 2003].



Experimental

The title compound was synthesized according to the literature method of Frederick & Liu (1999). Single crystals suitable for X-ray diffraction were obtained by slow evaporation of a solution in DMF–MeOH (1:1).

Crystal data	
C ₂₁ H ₁₅ NO	$D_x = 1.266 \text{ Mg m}^{-3}$
$M_r = 297.34$	Mo $K\alpha$ radiation
Monoclinic, Cc	Cell parameters from 1147
a = 36.589(5) Å	reflections
b = 5.3932 (7) Å	$\theta = 2.2 - 23.6^{\circ}$
c = 7.9572 (11) Å	$\mu = 0.08 \text{ mm}^{-1}$
$\beta = 96.689 \ (2)^{\circ}$	T = 292 (2) K
V = 1559.5 (4) Å ³	Block, colourless
Z = 4	$0.30\times0.20\times0.10$ mm
Data collection	
Bruker SMART CCD area-detector	1690 independent reflections
diffractometer	1360 reflections with $I > 2\sigma(I)$
φ and ω scans	$R_{\rm int} = 0.030$
Absorption correction: multi-scan	$\theta_{\rm max} = 27.0^{\circ}$
(SADABS; Sheldrick, 1996)	$h = -35 \rightarrow 46$
$T_{\min} = 0.977, \ T_{\max} = 0.992$	$k = -6 \rightarrow 6$
4343 measured reflections	$l = -10 \rightarrow 8$

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Refinement on F^2 $P[F^2 > 2\sigma(F^2)] = 0.045$	H-atom parameters constrained $w = 1/[\sigma^2(F^2) + (0.0786P)^2]$		
R[F > 20(F)] = 0.045 $wR(F^2) = 0.133$	where $P = (F_o^2 + 2F_c^2)/3$		
S = 1.08	$(\Delta/\sigma)_{\rm max} < 0.001$		
1690 reflections	$\Delta \rho_{\rm max} = 0.21 \text{ e A}^{\circ}$		
208 parameters	$\Delta \rho_{\rm min} = -0.24 \ {\rm e} \ {\rm A}^{-3}$		

Table 1

Hydrogen-bond geometry (Å, °).

$D=11\cdots A$	D-H	$\mathbf{H} \cdots \mathbf{A}$	$D \cdots A$	$D - H \cdots A$
$N1 - H1 \cdots O1^i$	0.86	2.42	3.250 (4)	163

Symmetry code: (i) x, y + 1, z.

After their location on a difference Fourier map, H atoms were placed in calculated positions and refined as riding with C–H = 0.93–0.97 Å, N–H = 0.86 Å, and U_{iso} = 1.2 or 1.5 times U_{eq} of the parent atom. Owing to the absence of any significant anomalous scatterers in the molecule, the Friedel pairs were merged before the final refinement.

Data collection: *SMART* (Bruker, 1997); cell refinement: *SAINT* (Bruker, 1999); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 2001); software used to prepare material for publication: *SHELXTL*.

The authors are grateful to the Central China Normal University, the National Natural Science Foundation of China (No.20472022), and the Hubei Province Natural Science Fund (Nos. 2004ABA085 and 2004ABC002) for financial support.

References

Allen, F. H., Kennard, O., Watson, D. G., Brammer, L., Orpen, A. G. & Taylor, R. (1987). J. Chem. Soc. Perkin Trans. 2, pp. S1–19.









Figure 2

The crystal packing, viewed approximately along the c axis. Intermolecular N-H···O hydrogen bonds are shown as dashed lines.

- Bruker (1997). SMART. Version 5.054. Bruker AXS Inc., Madison, Wisconsin, USA.
- Bruker (1999). SAINT. Version 6.01. Bruker AXS Inc., Madison, Wisconsin, USA.
- Bruker (2001). SHELXTL. Version 6.12. Bruker AXS Inc., Madison, Wisconsin, USA.
- Frederick, D. L. & Liu, W. (1999). J. Phys. Chem. A, 103, 9678-9686.
- Bowes, K. F., Glidewell, C., Low, J. N., Skakle, J. M. S. & Wardell, J. L. (2003). Acta Cryst. C59, 01–03.
- Sheldrick, G. M. (1996). SADABS. University of Göttingen, Germany.
- Sheldrick, G. M. (1997). SHELXS97 and SHELXL97. University of Göttingen, Germany.