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4-(Prop-2-yn-1-yloxy)benzene-1,2dicarbonitrile

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Key indicators: single-crystal X-ray study; T = 100 K; mean σ (C–C) = 0.003 Å; disorder in main residue; R factor = 0.059; wR factor = 0.163; data-to-parameter ratio = 13.0.

In the title compound, $C_{11}H_6N_2O$, the complete molecule is generated by the application of crystallographic twofold symmetry (the molecule is disordered about this axis). The prop-2-yn-1-yl residue is slightly twisted out of the plane of the benzene ring $[C-O-C-C \text{ torsion angle} = 173.1 (3)^{\circ}]$ and is orientated away from the nitrile substituents. In the crystal, supramolecular chains along the *a* axis, arising from $C-H \cdots N$ interactions, are connected into stacks along the c axis by $\pi - \pi$ interactions between the benzene rings [centroid-centroid distance = 3.6978 (6) Å = length of the *c* axis].

Related literature

For the solubilization and some applications of phthanocyanine dyes, see: Jiang et al. (2011); Sleven et al. (2001). For the synthesis of substituted phthalonitriles, see: Wöhrle et al. (1993); Wu et al. (1998); Li & Lieberman (2001); Sleven et al. (2001); Li et al. (2008); Seven et al. (2009); Foo et al. (2012).



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 $V = 942.62 (18) \text{ Å}^3$

Mo Ka radiation $\mu = 0.09 \text{ mm}^-$

 $0.15 \times 0.05 \times 0.05 \ \mathrm{mm}$

3258 measured reflections

1114 independent reflections

840 reflections with $I > 2\sigma(I)$

H atoms treated by a mixture of

independent and constrained

Z = 4

T = 100 K

 $R_{\rm int} = 0.038$

refinement

 $\Delta \rho_{\text{max}} = 0.41 \text{ e } \text{\AA}^{-3}$ $\Delta \rho_{\rm min} = -0.28 \text{ e } \text{\AA}^{-3}$

Experimental

Crystal data

$C_{11}H_6N_2O$	
$M_r = 182.18$	
Monoclinic, $C2/m$	
a = 11.4809 (9) Å	
b = 22.2091 (16) Å	
c = 3.6978 (6) Å	
$\beta = 91.304 \ (10)^{\circ}$	

Data collection

Agilent SuperNova Dual
diffractometer with an Atlas
detector
Absorption correction: multi-scan

(CrysAlis PRO; Agilent, 2012) $T_{\min} = 0.792, T_{\max} = 1.000$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.059$
$wR(F^2) = 0.163$
S = 1.08
1114 reflections
86 parameters
12 restraints

Table 1

Hydrogen-bond geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$\overline{C3-H3\cdots N1^{i}}$	0.95	2.62	3.509 (3)	155
Symmetry code: (i)	$-x + \frac{1}{2}, -y + \frac{1}{2},$	-z + 1.		

Data collection: CrysAlis PRO (Agilent, 2012); cell refinement: CrysAlis PRO; data reduction: CrysAlis PRO; program(s) used to solve structure: SHELXS97 (Sheldrick, 2008); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: ORTEP-3 (Farrugia, 1997) and DIAMOND (Brandenburg, 2006); software used to prepare material for publication: *publCIF* (Westrip, 2010).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HB6862).

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supplementary materials

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4-(Prop-2-yn-1-yloxy)benzene-1,2-dicarbonitrile

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Comment

Phthalocyanine dyes can be made soluble in water and organic solvents by addition of suitable alkoxy or aryloxy groups (Jiang *et al.*, 2011; Sleven *et al.*, 2001). This is most easily achieved from the correspondingly substituted phthalonitriles which, in turn, are either prepared by Sandmeyer reaction of alkyl or alkoxy functionalized dihalobenzenes (Li & Lieberman, 2001; Sleven *et al.*, 2001) or aryloxy / alkoxy displacement of the corresponding halophthalonitrile (Wöhrle *et al.*, 1993; Li *et al.*, 2008; Foo *et al.*, 2012) or 4-nitrophthalonitrile (Wu *et al.*, 1998; Seven *et al.*, 2009). The latter method is most suitable for preparing 4-alkoxyphthalonitriles and was used for preparing the title compound, 4-(prop-2-ylnyloxy)phthalonitrile (I).

In (I), Fig. 1, the complete molecule is generated by the application of 2-fold symmetry; the molecule is disordered about this axis. The O1 and C1 atoms lie -0.067 (3) and 0.059 (2) Å out of the plane through the benzene ring, respectively. The prop-2-yn-1-yl is twisted out of the plane of the benzene ring as seen in the value of the C4—O1—C5 —C6 torsion angle of 173.1 (3)° and is orientated in the opposite direction to the nitrile substituents.

In the crystal packing, supramolecular chains along the *a* axis feature owing to C—H…N interactions, Table 1, and 10membered {…HC₃N}₂ synthons, Fig. 2. Chains are connected into stacks along the *c* axis by π — π interactions between the benzene rings [inter-centroid distance = 3.6978 (6) Å = length of the *c* axis]. The layers inter-digitate along the *b* axis with no specific intermolecular interactions between them.

Experimental

The title compound was prepared by modification of literature procedures (Wu *et al.*, 1998; Seven *et al.*, 2009). Under a nitrogen atmosphere, anhydrous potassium carbonate (1.12 g, 8.1 mmol) was added in two portions at 1 h intervals to a solution of propargyl alcohol (1.5 ml, 26.0 mmol) and 4-nitrophthalonitrile (0.70 g, 4.04 mmol) in dry *N*,*N*-dimethyl-formamide (7 ml). After 96 h, the crude reaction mixture was poured into water (140 ml). The green precipitate was collected by vacuum filtration, washed with water and dried. The crude product was purified by silica gel column chromatography using dichloromethane as eluent to provide 0.4 g (63.9%) of a faintly coloured solid that was recrystallized from CH_2Cl_2 / hexane as colourless prisms. Melting point = 383 K. IR *v*/cm⁻¹: 3287, 3119, 3077, 2231, 2135, 1596, 1494, 1321, 1260. ¹H NMR 400 MHz (CDCl₃) δ : 7.75 (1*H*, d), 7.35 (1*H*, s), 7.28 (1*H*, d), 4.80 (2*H*, s), 2.62 (1*H*, s).

Refinement

With the exception of the acetylenic H-atom which was refined freely, carbon-bound H-atoms were placed in calculated positions [C—H = 0.95–0.99 Å, $U_{iso}(H) = 1.2U_{eq}(C)$] and were included in the refinement in the riding model approximation. The molecule is disordered over a 2-fold rotation axis in an exact 1:1 ratio. The anisotropic displacement

parameters of the O1 and C4 atoms were tightly restrained to be nearly isotropic.

Computing details

Data collection: *CrysAlis PRO* (Agilent, 2012); cell refinement: *CrysAlis PRO* (Agilent, 2012); data reduction: *CrysAlis PRO* (Agilent, 2012); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3* (Farrugia, 1997) and *DIAMOND* (Brandenburg, 2006); software used to prepare material for publication: *publCIF* (Westrip, 2010).



Figure 1

The molecular structure of (I) showing displacement ellipsoids at the 50% probability level. The molecule is disordered about the 2-fold axis - only one orientation is shown.



Figure 2

A view of the supramolecular chain along the *a* axis in (I). The C—H…N and interactions are shown as blue dashed lines.



Figure 3

A view of the supramolecular layer in the *ac* plane in (I). The C—H···O and π — π interactions are shown as blue and purple dashed lines, respectively.

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Crystal data

 $C_{11}H_6N_2O$ $M_r = 182.18$ Monoclinic, *C2/m*Hall symbol: -C 2y *a* = 11.4809 (9) Å *b* = 22.2091 (16) Å *c* = 3.6978 (6) Å *β* = 91.304 (10)° *V* = 942.62 (18) Å³ *Z* = 4

Data collection

Agilent SuperNova Dual	3258 measure
diffractometer with an Atlas detector	1114 independ
Radiation source: fine-focus sealed tube	840 reflection
Mirror monochromator	$R_{\rm int} = 0.038$
Detector resolution: 10.4041 pixels mm ⁻¹	$\theta_{\rm max} = 27.6^{\circ}, \ell$
ω scan	$h = -14 \rightarrow 14$
Absorption correction: multi-scan	$k = -26 \rightarrow 28$
(CrysAlis PRO; Agilent, 2012)	$l = -4 \rightarrow 4$
$T_{\min} = 0.792, \ T_{\max} = 1.000$	

Refinement

Refinement on F^2 Least-squares matrix: full $R[F^2 > 2\sigma(F^2)] = 0.059$ $wR(F^2) = 0.163$ S = 1.081114 reflections 86 parameters 12 restraints Primary atom site location: structure-invariant direct methods F(000) = 376 $D_x = 1.284 \text{ Mg m}^{-3}$ Mo K\alpha radiation, $\lambda = 0.71073 \text{ Å}$ Cell parameters from 938 reflections $\theta = 3.3-27.5^{\circ}$ $\mu = 0.09 \text{ mm}^{-1}$ T = 100 KPrism, colourless $0.15 \times 0.05 \times 0.05 \text{ mm}$

3258 measured reflections 1114 independent reflections 840 reflections with $I > 2\sigma(I)$ $R_{int} = 0.038$ $\theta_{max} = 27.6^{\circ}, \ \theta_{min} = 3.3^{\circ}$ $h = -14 \rightarrow 14$ $k = -26 \rightarrow 28$ $l = -4 \rightarrow 4$

Secondary atom site location: difference Fourier map Hydrogen site location: inferred from neighbouring sites H atoms treated by a mixture of independent and constrained refinement $w = 1/[\sigma^2(F_o^2) + (0.0718P)^2 + 0.8054P]$ where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{max} < 0.001$ $\Delta\rho_{max} = 0.41$ e Å⁻³ $\Delta\rho_{min} = -0.28$ e Å⁻³

Special details

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

Refinement. Refinement of F^2 against ALL reflections. The weighted *R*-factor *wR* and goodness of fit *S* are based on F^2 , conventional *R*-factors *R* are based on *F*, with *F* set to zero for negative F^2 . The threshold expression of $F^2 > \sigma(F^2)$ is used only for calculating *R*-factors(gt) *etc.* and is not relevant to the choice of reflections for refinement. *R*-factors based on F^2 are statistically about twice as large as those based on *F*, and *R*- factors based on ALL data will be even larger.

	x	У	Ζ	$U_{ m iso}$ */ $U_{ m eq}$	Occ. (<1)
01	0.1308 (2)	0.45076 (11)	0.1768 (8)	0.0273 (7)	0.50
N1	0.15188 (16)	0.19174 (9)	0.2675 (6)	0.0376 (6)	
C1	0.10982 (16)	0.23658 (9)	0.1829 (6)	0.0260 (5)	
C2	0.05529 (16)	0.29272 (8)	0.0857 (5)	0.0219 (5)	
C3	0.1103 (2)	0.34676 (10)	0.1663 (6)	0.0349 (6)	
H3	0.1853	0.3469	0.2801	0.042*	
C4	0.0555 (3)	0.40028 (10)	0.0803 (7)	0.0446 (7)	
H4	0.0936	0.4374	0.1309	0.054*	0.50
C5	0.2499 (3)	0.44609 (17)	0.3108 (13)	0.0274 (10)	0.50
H5A	0.2962	0.4211	0.1458	0.033*	0.50
H5B	0.2522	0.4272	0.5535	0.033*	0.50
C6	0.2980 (3)	0.5082 (7)	0.3303 (12)	0.030 (3)	0.50
C7	0.3440 (4)	0.5544 (2)	0.3459 (17)	0.0463 (14)	0.50
H7	0.386 (5)	0.595 (3)	0.363 (16)	0.054 (16)*	0.50

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (A^2)

Atomic displacement parameters $(Å^2)$

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
01	0.0258 (13)	0.0187 (12)	0.0370 (15)	0.0001 (10)	-0.0069 (11)	-0.0008 (11)
N1	0.0332 (10)	0.0410 (11)	0.0383 (12)	0.0126 (9)	-0.0029 (8)	0.0074 (9)
C1	0.0191 (9)	0.0345 (12)	0.0242 (11)	-0.0003 (8)	-0.0019 (8)	0.0004 (8)
C2	0.0206 (10)	0.0234 (10)	0.0217 (10)	-0.0009 (7)	-0.0001 (8)	-0.0007 (7)
C3	0.0401 (12)	0.0399 (13)	0.0251 (12)	-0.0182 (10)	0.0071 (9)	-0.0077 (9)
C4	0.0728 (15)	0.0262 (10)	0.0355 (13)	-0.0161 (10)	0.0172 (11)	-0.0063 (9)
C5	0.0220 (19)	0.0167 (19)	0.043 (3)	0.0005 (17)	-0.0104 (17)	-0.0001 (16)
C6	0.0262 (16)	0.013 (9)	0.051 (2)	-0.004 (2)	-0.0130 (15)	-0.001 (2)
C7	0.033 (3)	0.027 (2)	0.078 (4)	0.001 (2)	-0.019 (2)	-0.001(2)

Geometric parameters (Å, °)

01—C5	1.447 (5)	C4—C4 ⁱ	1.393 (6)
O1—C4	1.455 (3)	C4—H4	0.9500
N1-C1	1.147 (3)	C5—C6	1.487 (14)
C1—C2	1.437 (3)	C5—H5A	0.9900
С2—С3	1.385 (3)	C5—H5B	0.9900
$C2-C2^i$	1.406 (4)	C6—C7	1.155 (16)
C3—C4	1.379 (3)	С7—Н7	1.02 (6)

С3—Н3	0.9500		
C5—O1—C4	125.5 (3)	С3—С4—Н4	119.8
N1—C1—C2	178.4 (2)	$C4^{i}$ — $C4$ — $H4$	119.8
$C3-C2-C2^{i}$	119.96 (13)	O1—C5—C6	107.3 (3)
C3—C2—C1	120.26 (18)	O1—C5—H5A	110.3
$C2^{i}$ — $C2$ — $C1$	119.77 (10)	C6—C5—H5A	110.3
C4—C3—C2	119.6 (2)	O1—C5—H5B	110.3
С4—С3—Н3	120.2	C6—C5—H5B	110.3
С2—С3—Н3	120.2	H5A—C5—H5B	108.5
$C3-C4-C4^{i}$	120.43 (14)	C7—C6—C5	174.6 (7)
C3—C4—O1	110.0 (2)	С6—С7—Н7	178 (3)
C4 ⁱ —C4—O1	129.56 (15)		
C2 ⁱ —C2—C3—C4	0.6 (4)	C5—O1—C4—C3	5.5 (5)
C1—C2—C3—C4	-178.2 (2)	C5	-173.5 (4)
$C2-C3-C4-C4^{i}$	1.2 (4)	C4—O1—C5—C6	173.1 (3)
C2—C3—C4—O1	-177.9 (2)		

Symmetry code: (i) -x, y, -z.

Hydrogen-bond geometry (Å, °)

D—H···A	<i>D</i> —Н	H···A	$D \cdots A$	D—H···A
C3—H3…N1 ⁱⁱ	0.95	2.62	3.509 (3)	155

Symmetry code: (ii) -x+1/2, -y+1/2, -z+1.