

Ufuk Çoruh,^{a*} Nesuhi Akdemir,^a
Erbil Açar,^a Ezequiel M.
Vázquez-López^b and Ahmet
Erdönmez^a

^aOndokuz Mayıs University, Art and Science Faculty, Department of Physics, 55139 Samsun, Turkey, and ^bDepartamento de Química Inorgánica, Facultad de Ciencias-Química, Universidade de Vigo, 36200-Vigo, Galicia, Spain

Correspondence e-mail: ucoruh@hotmail.com

Key indicators

Single-crystal X-ray study

$T = 293\text{ K}$

Mean $\sigma(\text{C}-\text{C}) = 0.007\text{ \AA}$

R factor = 0.046

wR factor = 0.154

Data-to-parameter ratio = 17.5

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

4,4'-[2,2'-(Piperidine-1,4-diyl-diethylene)di(tosylimino)]diphthalonitrile

The title compound, $\text{C}_{38}\text{H}_{36}\text{N}_8\text{O}_4\text{S}_2$, has a center of symmetry and the central six-membered ring shows a chair conformation. The dihedral angle between the ring carrying the tosyl group and the phthalonitrile ring is $36.98(11)^\circ$. The crystal structure is stabilized by an intramolecular $\text{C}-\text{H}\cdots\text{N}$ and two intermolecular $\text{C}-\text{H}\cdots\text{O}$ close contacts.

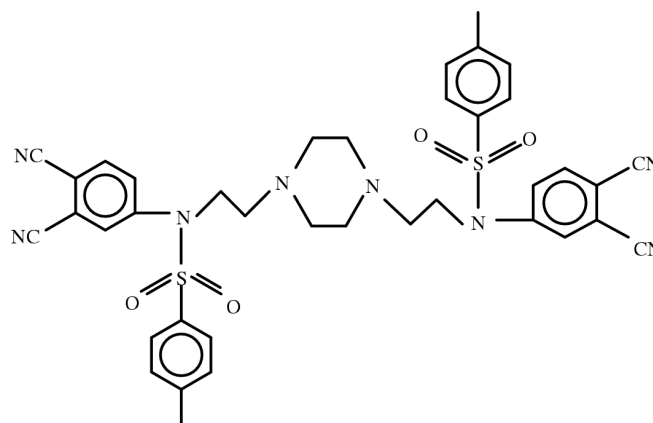
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Comment

The title compound, (I), is a precursor in the synthesis of network phthalocyanine polymers (Açar *et al.*, 1995). For many years, phthalocyanines have attracted continued interest in various research fields, such as chemical sensors, electrochromism, batteries, photodynamic therapy, semiconductive materials, liquid crystals and non-linear optics (Leznoff & Lever, 1989–1996; McKeown, 1998).



(I)

The atom-numbering scheme and a displacement ellipsoid plot of (I) are shown in Fig. 1. The geometry of the S atoms is distorted from the tetrahedral configuration, with the largest deviation in the $\text{O}-\text{S}-\text{O}$ angle [$\text{O}1-\text{S}1-\text{O}2 = 120.41(18)^\circ$] and the angles $\text{O}1-\text{S}1-\text{N}3$ and $\text{O}2-\text{S}1-\text{N}3$ [$107.37(18)$ and $106.16(18)^\circ$, respectively]. The $\text{S}1-\text{N}3$ distance is $1.628(3)\text{ \AA}$, very close to those in the literature (Işık *et al.* 1999) and in contrast with the previously observed range of $1.63\text{--}1.69\text{ \AA}$ (Kálmán *et al.*, 1981; Öztürk *et al.*, 2000). This shortening is due to the electron-withdrawing character of the diphthalonitrile group, as in 10,11-dibromo-3,6-ditosyl-3,6-diazabicyclo[6.4.0]dodeca-1(8),9,11-triene (Işık *et al.*, 1999).

The $\text{S}1-\text{C}13$ bond distance and average $\text{S}=\text{O}$ bond length are $1.765(4)$ and $1.433(3)\text{ \AA}$, respectively. The bond lengths in the literature (Öztürk *et al.*, 2000) are $1.742(6)$ and $1.430(5)\text{ \AA}$, respectively. The $\text{C}1\equiv\text{N}1$ and $\text{C}2\equiv\text{N}2$ bond distances are $1.130(5)$ and $1.142(5)\text{ \AA}$, respectively, and show $\text{N}\equiv\text{C}$ triple-bond character. The ring composed of atoms $\text{C}18$,

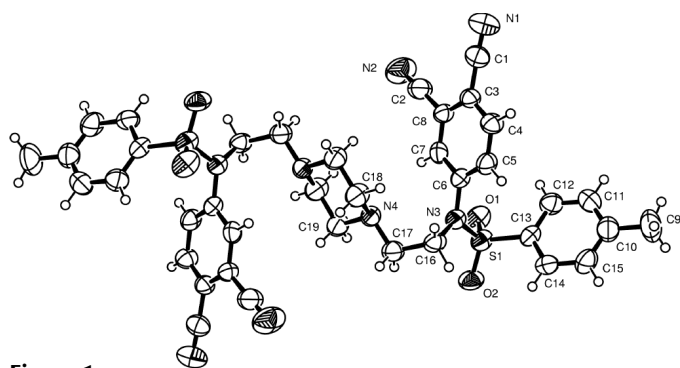


Figure 1

A view of (I) with the atomic numbering scheme. Displacement ellipsoids are drawn at the 50% probability level.

C19 and N4 and their symmetry equivalents is almost in a chair conformation. The crystal structure of (I) is stabilized by the intramolecular C7—H7···N4 interaction, and the intermolecular C15—H15···O1ⁱ and C18—H18A···O2ⁱⁱ close contacts (see Table 2 for details and symmetry code).

Experimental

1,4-Bis[*N*-(2-tosylaminoethyl)]-1,4-diazacyclohexane (13.16 g, 27.42 mmol; Hancock *et al.*, 1990) was dissolved in dry dimethyl sulfoxide (150 ml) under nitrogen and 4-nitrophthalonitrile (10.11 g, 58.43 mmol) was added. After stirring for 30 min, finely ground anhydrous K₂CO₃ (19.73 g, 142.97 mmol) was added portionwise over a period of 2 h with efficient stirring. The reaction mixture was stirred under nitrogen at room temperature for 24 h. Water was then added and the product filtered off and washed with water until the filtrate was neutral. The product was then refluxed in methanol, filtered and dried. The title compound, (I), was crystallized from dimethylformamide *via* slow evaporation at room temperature (yield 8.13 g, 40.51%). Analysis calculated for C₃₈H₃₆N₈O₄S₂: C 62.28, H 4.95, N 15.29%; found: C 62.65, H 4.93, N 15.44%. Full IR, ¹H and ¹³C NMR spectral data have been deposited elsewhere (Çelebi, 2002).

Crystal data

C ₃₈ H ₃₆ N ₈ O ₄ S ₂	$D_x = 1.295 \text{ Mg m}^{-3}$
$M_r = 732.89$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/c$	Cell parameters from 9948 reflections
$a = 8.4238 (11) \text{ \AA}$	$\theta = 1.4\text{--}28.0^\circ$
$b = 29.217 (4) \text{ \AA}$	$\mu = 0.19 \text{ mm}^{-1}$
$c = 7.7120 (11) \text{ \AA}$	$T = 293 (2) \text{ K}$
$\beta = 98.052 (3)^\circ$	Prism, colourless
$V = 1879.4 (5) \text{ \AA}^3$	$0.21 \times 0.19 \times 0.06 \text{ mm}$
$Z = 2$	

Data collection

Bruker CCD area-detector diffractometer	1241 reflections with $I > 2\sigma(I)$
φ and ω scans	$R_{\text{int}} = 0.078$
Absorption correction: none	$\theta_{\text{max}} = 28.0^\circ$
9948 measured reflections	$h = -10 \rightarrow 11$
4124 independent reflections	$k = -33 \rightarrow 38$
	$l = -10 \rightarrow 9$

Refinement

Refinement on F^2	H-atom parameters constrained
$R[F^2 > 2\sigma(F^2)] = 0.046$	$w = 1/[\sigma^2(F_o^2) + (0.0613P)^2]$
$wR(F^2) = 0.154$	where $P = (F_o^2 + 2F_c^2)/3$
$S = 0.72$	$(\Delta/\sigma)_{\text{max}} < 0.001$
4124 reflections	$\Delta\rho_{\text{max}} = 0.22 \text{ e \AA}^{-3}$
235 parameters	$\Delta\rho_{\text{min}} = -0.19 \text{ e \AA}^{-3}$

Table 1

Selected geometric parameters (\AA , $^\circ$).

N3—C6	1.440 (5)	C10—C9	1.499 (6)
N3—C16	1.465 (5)	N4—C19	1.453 (4)
N3—S1	1.628 (3)	N4—C17	1.456 (4)
S1—O1	1.429 (3)	N4—C18	1.456 (5)
S1—O2	1.438 (3)	C1—N1	1.130 (5)
S1—C13	1.765 (4)	N2—C2	1.142 (5)
C16—C17	1.523 (5)		
C6—N3—C16	118.3 (3)	O2—S1—N3	106.16 (18)
C6—N3—S1	119.0 (3)	O1—S1—C13	106.8 (2)
C16—N3—S1	120.8 (3)	O2—S1—C13	109.1 (2)
O1—S1—O2	120.41 (18)	N3—S1—C13	106.16 (17)
O1—S1—N3	107.37 (18)		

Table 2

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

$D\text{---}H\cdots A$	$D\text{---}H$	$H\cdots A$	$D\cdots A$	$D\text{---}H\cdots A$
C7—H7···N4	0.93	2.95	3.319 (5)	105
C15—H15···O1 ⁱ	0.93	2.69	3.559 (5)	155
C18—H18A···O2 ⁱⁱ	0.97	2.60	3.248 (5)	123

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

All H atoms were positioned geometrically and refined using a riding model, fixing the aromatic C—H distance at 0.93 \AA , the methylene C—H distance at 0.97 \AA and methyl-group C—H distance at 0.96 \AA .

Data collection: SMART (Bruker, 1998); cell refinement: SAINT (Bruker, 1998); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEP-3 (Farrugia, 1997) and PLATON (Spek, 1990); software used to prepare material for publication: SHELXL97 and PARST (Nardelli, 1995).

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