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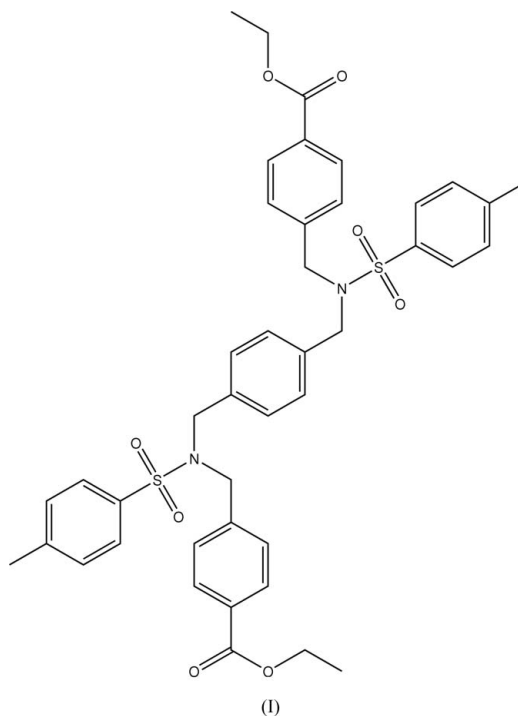
Key indicators

Single-crystal X-ray study
T = 123 K
Mean $\sigma(\text{C}-\text{C}) = 0.002 \text{ \AA}$
R factor = 0.033
wR factor = 0.093
Data-to-parameter ratio = 14.6For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.1,4-Bis{N-[4-(ethoxycarbonyl)benzyl]-N-(4-tolyl-
sulfonyl)aminomethyl}benzene

The title compound, $\text{C}_{42}\text{H}_{44}\text{N}_2\text{O}_8\text{S}_2$, features a molecular thread. The molecules have crystallographic C_i symmetry. The crystal packing shows staples stabilized by weak $\text{C}-\text{H}\cdots\text{O}$ interactions.

Comment

In supramolecular chemistry, and especially nanochemistry, macrocyclic and concave hydrocarbons play an important role. Nanometric scale molecules are fascinating due to their often appealing architecture, high symmetry and host-guest interactions. Linear molecular threads play an important role in the synthesis of nanometer-scaled molecular ribbons and belts (Schwierz & Vögtle, 1999).



The title compound, (I), features a molecular thread. The backbone of the molecules (disregarding the tosyl residues) shows the form of the letter S. The molecules have crystallographic C_i symmetry (Fig. 1) and form, similar molecular ribbons (Breidenbach *et al.*, 1995), staples (Figs. 2 and 3) which are stabilized by weak $\text{C}-\text{H}\cdots\text{O}$ interactions (Table 1) (Steiner, 2002).

Experimental

The title compound was synthesized by the reaction of two equivalents of ethyl 4-(bromomethyl)benzoate and one equivalent of 1,4-

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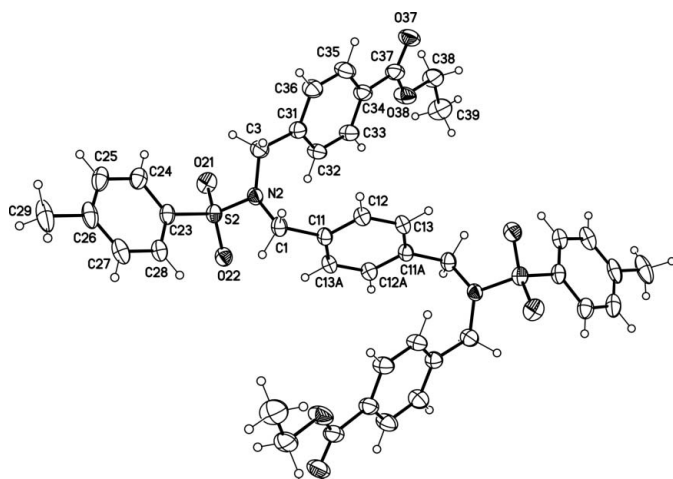


Figure 1
View of the molecule of the title compound showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 50% probability level. H atoms are represented by circles of arbitrary size. [Symmetry code: (A, unlabelled atoms): $1 - x, 1 - y, 1 - z$.]

bis[*N*-(4-tolylsulfonyl)aminomethyl]benzene in *N,N*-dimethylformamide and K_2CO_3 . The precipitate was recrystallized from CH_2Cl_2 (Schwierz, 1999).

Crystal data

$C_{42}H_{44}N_2O_8S_2$	$Z = 1$
$M_r = 768.91$	$D_x = 1.320 \text{ Mg m}^{-3}$
Triclinic, $P\bar{1}$	Mo $K\alpha$ radiation
$a = 6.7007 (2) \text{ \AA}$	Cell parameters from 15012 reflections
$b = 11.7108 (5) \text{ \AA}$	$\theta = 3.2\text{--}28.3^\circ$
$c = 13.1888 (5) \text{ \AA}$	$\mu = 0.19 \text{ mm}^{-1}$
$\alpha = 78.912 (2)^\circ$	$T = 123 (2) \text{ K}$
$\beta = 80.801 (2)^\circ$	Plates, colourless
$\gamma = 73.492 (2)^\circ$	$0.35 \times 0.30 \times 0.05 \text{ mm}$
$V = 967.64 (6) \text{ \AA}^3$	

Data collection

Nonius KappaCCD diffractometer	$R_{\text{int}} = 0.021$
φ scans	$\theta_{\text{max}} = 28.3^\circ$
Absorption correction: none	$h = -8 \rightarrow 8$
15012 measured reflections	$k = -14 \rightarrow 14$
3582 independent reflections	$l = -15 \rightarrow 15$
3145 reflections with $I > 2\sigma(I)$	

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0481P)^2 + 0.2922P]$
$R[F^2 > 2\sigma(F^2)] = 0.033$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.093$	$(\Delta/\sigma)_{\text{max}} = 0.001$
$S = 1.04$	$\Delta\rho_{\text{max}} = 0.24 \text{ e \AA}^{-3}$
3582 reflections	$\Delta\rho_{\text{min}} = -0.38 \text{ e \AA}^{-3}$
245 parameters	
H-atom parameters constrained	

Table 1

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

$D\text{---}H\cdots A$	$D\text{---}H$	$H\cdots A$	$D\cdots A$	$D\text{---}H\cdots A$
$C28\text{---}H28\cdots O37^i$	0.95	2.53	3.3341 (19)	143

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

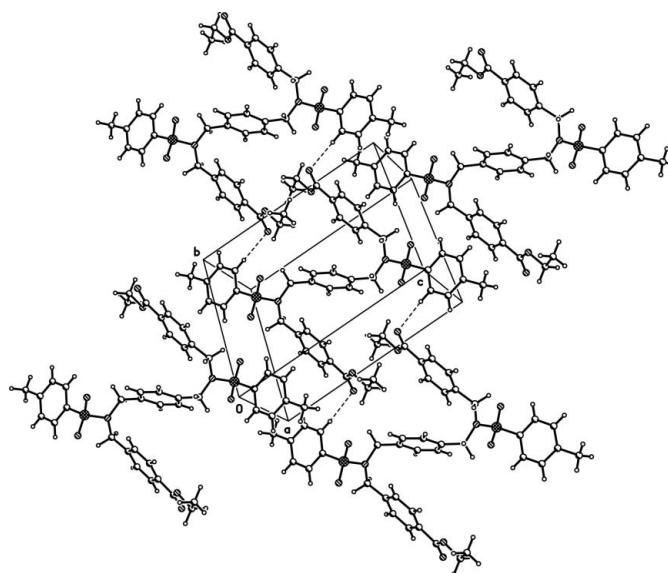


Figure 2
The crystal packing of the title compound. Hydrogen bonds are shown as dashed lines.

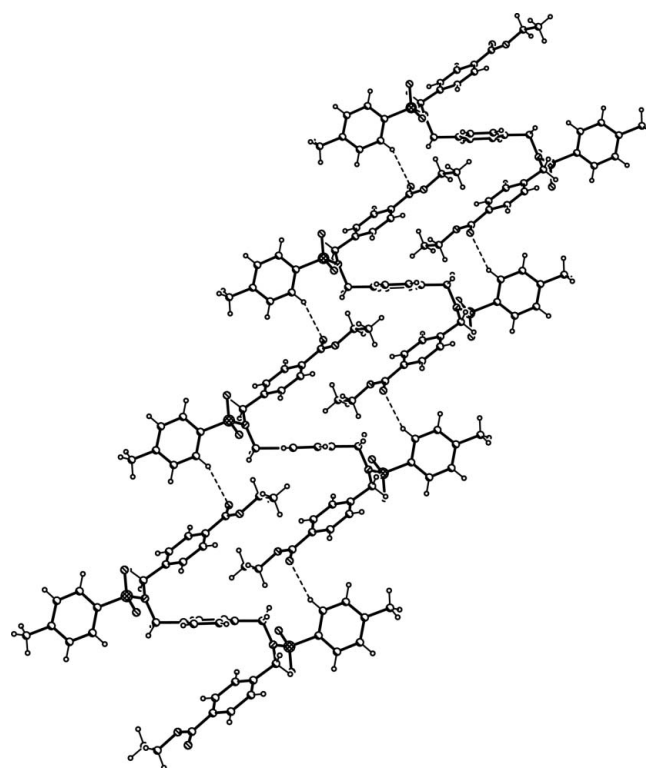


Figure 3
The staple formation of the title compound. Hydrogen bonds are shown as dashed lines.

All H atoms were initially located in a difference Fourier map. The methyl H atoms were then constrained to an ideal geometry, with C—H distances of 0.98 \AA and with $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{C})$. All other H atoms were placed in geometrically idealized positions and constrained to ride on their parent atoms, with C—H distances in the range $0.95\text{--}0.99 \text{ \AA}$ and with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$.

Data collection: *COLLECT* (Nonius, 1997); cell refinement: *DENZO-SMN* (Otwinowski & Minor, 1997); data reduction:

DENZO-SMN; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1990); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Sheldrick, 2001); software used to prepare material for publication: *SHELXL97*.

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